GenCore version 4.5

OM nucleic - nucleic search, using sw model

Run on: December 8, 2001, 19:38:26 ; Search time 70.08 Seconds
(without alignments)
12804.022 Million cell updates/sec

Title: US-08-153-397A-1
Perfect score: 3962
Sequence: 1 CGGGCCTGAGACTGGGGTGA.....AAAAAAAAAAACCGGAATTC 3962

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 351203 seqs, 113238999 residues

Total number of hits satisfying chosen parameters: 702406

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued_Patents_NA:*
1: /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/2/ina/5B_COMB.seq:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Result No.	Score	% Query Match	Length	DB	ID	Description
	1	3962	100.0	3962	1	US-08-336-343A-1	Sequence 1, Appli
	2	3451	87.1	3637	1	US-08-445-640-3	Sequence 3, Appli
	3	3451	87.1	3637	3	US-08-170-558-3	Sequence 3, Appli
	4	3451	87.1	3637	3	US-08-447-314-3	Sequence 3, Appli
	5	3451	87.1	3637	3	US-08-445-461-3	Sequence 3, Appli
	6	1192.2	30.1	1197	1	US-08-445-640-7	Sequence 7, Appli
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c	11	642	16.2	3157	1	US-08-336-343A-5	Sequence 5, Appli
	12	639.8	16.1	3120	1	US-08-456-647B-19	Sequence 19, Appl
	13	639.8	16.1	3120	2	US-08-237-401A-19	Sequence 19, Appl
	14	182.2	4.6	2820	1	US-08-286-305A-4	Sequence 4, Appli
	15	182.2	4.6	2820	2	US-08-441-104A-4	Sequence 4, Appli
	16	182.2	4.6	2820	2	US-08-440-816A-4	Sequence 4, Appli
	17	182.2	4.6	2820	4	US-09-417-381A-4	Sequence 4, Appli
	18	180.6	4.6	2301	1	US-08-306-691B-23	Sequence 23, Appl
	19	180.6	4.6	2301	5	PCT-US93-06251-78	Sequence 78, Appl
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	27	180.6	4.6	3194	3	US-08-444-622A-1	Sequence 1, Appli
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	38	158	4.0	4092	2	US-08-469-537A-106	Sequence 106, App
	39	147.2	3.7	3398	5	PCT-US95-08493-12	Sequence 12, Appl
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	41	139	3.5	2580	5	PCT-US95-08493-18	Sequence 18, Appl
	42	139	3.5	2604	5	PCT-US95-08493-20	Sequence 20, Appl
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	45	130.4	3.3	2869	1	US-08-374-834-2	Sequence 2, Appli

ALIGNMENTS

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RESULT 1
US-08-336-343A-1
; Sequence 1, Application US/08336343A
; Patent No. 5677144
; GENERAL INFORMATION:
;   APPLICANT: Ullrich, Axel
;   APPLICANT: Alves, Frauke
;   TITLE OF INVENTION: CCK-2, A No. 5677144el Receptor Tyrosine Kinase
;   NUMBER OF SEQUENCES: 43
;   CORRESPONDENCE ADDRESS:
;     ADDRESSEE: Pennie & Edmonds
;     STREET: 1155 Avenue of the Americas
;     CITY: New York
;     STATE: New York
;     COUNTRY: U.S.A.
;     ZIP: 10036-2711
;   COMPUTER READABLE FORM:
;     MEDIUM TYPE: Floppy disk
;     COMPUTER: IBM PC compatible
;     OPERATING SYSTEM: PC-DOS/MS-DOS
;     SOFTWARE: PatentIn Release #1.0, Version #1.30
;   CURRENT APPLICATION DATA:
;     APPLICATION NUMBER: US/08/336,343A
;     FILING DATE: 08-NOV-1994
;     CLASSIFICATION: 435
;   ATTORNEY/AGENT INFORMATION:
;     NAME: Coruzzi, Laura A.
;     REGISTRATION NUMBER: 30,742
;     REFERENCE/DOCKET NUMBER: 7683-065
;   TELECOMMUNICATION INFORMATION:
;     TELEPHONE: (212) 790-9090
;     TELEFAX: (212) 869-9741/8864
;     TELEX: 66141 PENNIE
; INFORMATION FOR SEQ ID NO: 1:
;   SEQUENCE CHARACTERISTICS:
;     LENGTH: 3962 base pairs
;     TYPE: nucleic acid
;     STRANDEDNESS: double
;     TOPOLOGY: unknown
;   MOLECULE TYPE: cDNA
;   HYPOTHETICAL: NO
;   ANTI-SENSE: NO
;   FEATURE:
;     NAME/KEY: CDS
;     LOCATION: 321..3077
US-08-336-343A-1

  Query Match 100.0%; Score 3962; DB 1; Length 3962;
  Best Local Similarity 100.0%; Pred. No. 0;
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Matches 3962; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db     181 CCCGGGTCGGACGCCTGGGTCTGCCGGGAAGAGCGATGAGAGGTGTCTGAAGGTGGCTAT 240

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Db   2161 AGAAGCTTGGCGAGGGCCAGTTTGGGGAGGTGCACCTGTGTGAGGTCGACAGCCCTCAAG 2220

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Qy   2401 TGGGCGTGTGTGTGCAGGACGACCCCCTCTGCATGATTACTGACTACATGGAGAACGGCG 2460
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Qy   3901 GCACTAGGCAGGTAATAATAAAGGTTGAGTTTTCCACAAAAAAAAAAAAAAAAACCGGAAT 3960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3901 GCACTAGGCAGGTAATAATAAAGGTTGAGTTTTCCACAAAAAAAAAAAAAAAAACCGGAAT 3960

Qy   3961 TC 3962
          ||
Db   3961 TC 3962
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RESULT 2
US-08-445-640-3
; Sequence 3, Application US/08445640
; Patent No. 5709858
; GENERAL INFORMATION:
; APPLICANT: Godowski, Paul J.
; APPLICANT: Mark, Melanie R.
; APPLICANT: Scadden, David T.
; APPLICANT: Baker, Kevin P.
; APPLICANT: Baron, Will F.
; TITLE OF INVENTION: Protein Tyrosine Kinases
; NUMBER OF SEQUENCES: 35
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Genentech, Inc.
; STREET: 460 Point San Bruno Blvd
; CITY: South San Francisco
; STATE: California
; COUNTRY: USA
; ZIP: 94080
; COMPUTER READABLE FORM:
; MEDIUM TYPE: 5.25 inch, 360 Kb floppy disk

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;       COMPUTER:  IBM PC compatible
;       OPERATING SYSTEM:  PC-DOS/MS-DOS
;       SOFTWARE:  patin (Genentech)
;     CURRENT APPLICATION DATA:
;       APPLICATION NUMBER:  US/08/445,640
;       FILING DATE:  22-MAY-1995
;       CLASSIFICATION:  435
;     PRIOR APPLICATION DATA:
;       APPLICATION NUMBER:  08/170558
;       FILING DATE:  20-DEC-1993
;     PRIOR APPLICATION DATA:
;       APPLICATION NUMBER:  08/157563
;       FILING DATE:  23-NOV-1993
;     ATTORNEY/AGENT INFORMATION:
;       NAME:  Hasak, Janet E.
;       REGISTRATION NUMBER:  28,616
;       REFERENCE/DOCKET NUMBER:  834C2
;     TELECOMMUNICATION INFORMATION:
;       TELEPHONE:  415/225-1896
;       TELEFAX:  415/952-9881
;       TELEX:  910/371-7168
;   INFORMATION FOR SEQ ID NO:  3:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  3637 bases
;       TYPE:  nucleic acid
;       STRANDEDNESS:  single
;       TOPOLOGY:  linear
US-08-445-640-3


  Query Match             87.1%;  Score 3451;  DB 1;  Length 3637;
  Best Local Similarity   97.0%;  Pred. No. 0;
  Matches 3589;  Conservative    0;  Mismatches    5;  Indels  105;  Gaps    3;

Qy    256 GTTGGACTTGAAGGAATGCCAAGAGATGCTGCCCCCACCCCCTTAGGCCCGAGGGATCAG 315
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Db     17 GTTGGACTTGAAGGAATGCCAAGAGATGCTGCCCCCACCCCCTTAGGCCCGAGGGATCAG 76

Qy    316 GAGCTATGGGACCAGAGGCCCTGTCATCTTTACTGCTGCTGCTCTTGGTGGCAAGTGGAG 375
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     77 GAGCTATGGGACCAGAGGCCCTGTCATCTTTACTGCTGCTGCTCTTGGTGGCAAGTGGAG 136

Qy    376 ATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAGG 435
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Db    137 ATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAGG 196

Qy    436 ACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCCG 495
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    197 ACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCCG 256

Qy    496 CCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCGG 555
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    257 CCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCGG 316

Qy    556 TGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTCCACCTGGTGGCTC 615
          ||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||
Db    317 TGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTGCACCTGGTGGCTC 376

Qy    616 TGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTACC 675
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    377 TGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTACC 436

Qy    676 GGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAGG 735
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Db    437 GGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAGG 496

Qy    736 AGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCCA 795
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    497 AGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCCA 556

Qy    796 TGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGTGTCTGTCTGC 855
          ||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db    557 TGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGCGTCTGTCTGC 616

Qy    856 GGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTGG 915
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    617 GGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTGG 676

Qy    916 GGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACATA 975
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Db    677 GGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACATA 736

Qy    976 CCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGG 1035
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Db    737 CCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGG 796

Qy   1036 ATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGA 1095
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Db    797 ATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGA 856

Qy   1096 GCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGG 1155
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    857 GCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGG 916

Qy   1156 CCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCTG 1215
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    917 CCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCTG 976

Qy   1216 GCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGC 1275
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Db    977 GCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGC 1036

Qy   1276 GCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTG 1335
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Db    1037 GCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTG 1096

Qy   1336 GCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCT 1395
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Db    1097 GCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCT 1156

Qy   1396 TCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCA 1455
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1157 TCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCA 1216

Qy   1456 CCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGG 1515
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Db    1217 CCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGG 1276

Qy   1516 AGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCCATCC 1575
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Db    1277 AGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGCCCGAGGGGAGCCCGACCGCCATCC 1336

Qy   1576 TCATCGGCTGCCTGGTGGCCATCATCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCT 1635
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Db    1397 GGCGGCTGCACTGGCGCAGGCTCCTCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGC 1456

Qy   1696 TGACGGTTCACCTCTCTGTCCCTGGGGACACTATCCTCATCAACAACCGCCCAGGTCCTA 1755
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Db    1457 TGACGGTTCACCTCTCTGTCCCTGGGGACACTATCCTCATCAACAACCGCCCAGGTCCTA 1516

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Db    1517 GAGAGCCACCCCCGTACCAGGAGCCCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCT 1576

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Db    1577 GTGTCCCCAATGGCTCTGCGTTGCTGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCA 1636

Qy   1876 CTTACGCCCGTCCCCCTCGAGGCCCGGGCCCCCCCACACCCGCCTGGGCCAAACCCACCA 1935
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Db    1637 CTTACGCCCGTCCCCCTCGAGGCCCGGGCCCCCCCACACCCGCCTGGGCCAAACCCACCA 1696
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Qy   1936 ACACCCAGGCCTACAGTGGGGACTATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGC 1995
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Db   1697 ACACCCAGGCCTACAGTGGGGACTATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGC 1756

Qy   1996 CCCCACCTCCCCAGAACAGCGTCCCCCATTATGCCGAGGCTGACATTGTTACCCTGCAGG 2055
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Db   1757 CCCCACCTCCCCAGAACAGCGTCCCCCATTATGCCGAGGCTGACATTGTTACCCTGCAGG 1816

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Db   1817 GCGTCACCGGGGGCAACACCTATGCTGTGCCTGCACTGCCCCCAGGGGCAGTCGGGGATG 1876

Qy   2116 GGCCCCCCAGAGTGGATTTCCCTCGATCTCGACTCCGCTTCAAGGAGAAGCTTGGCGAGG 2175
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Qy   2176 GCCAGTTTGGGGAGGTGCACCTGTGTGAGGTCGACAGCCCTCAAGATCTGGTCAGTCTTG 2235
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Qy   2236 ATTTCCCCCTTAATGTGCGTAAGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGC 2295
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Qy   2416 AGGACGACCCCCTCTGCATGATTACTGACTACATGGAGAACGGCGACCTCAACCAGTTCC 2475
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Db   2159 AGGACGACCCCCTCTGCATGATTACTGACTACATGGAGAACGGCGACCTCAACCAGTTCC 2218

Qy   2476 TCAGTGCCCACCAGCTGGAGGACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTG 2535
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Db   2219 TCAGTGCCCACCAGCTGGAGGACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTG 2278

Qy   2536 CGCAGGGGCCCACCATCAGCTACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCG 2595
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Db   2279 CGCAGGGGCCCACCATCAGCTACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCG 2338

Qy   2596 GCATGCGCTATCTGGCCACACTCAACTTTGTACATCGGGACCTGGCCACGCGGAACTGCC 2655
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Db   2339 GCATGCGCTATCTGGCCACACTCAACTTTGTACATCGGGACCTGGCCACGCGGAACTGCC 2398

Qy   2656 TAGTTGGGGAAAATTTCACCATCAAAATCGCAGACTTTGGCATGAGCCGGAACCTCTATG 2715
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Qy   2716 CTGGGGACTATTACCGTGTGCAGGGCCGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGG 2775
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Db   2459 CTGGGGACTATTACCGTGTGCAGGGCCGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGG 2518

Qy   2776 AGTGCATCCTCATGGGGAAGTTCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCC 2835
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2519 AGTGCATCCTCATGGGGAAGTTCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCC 2578

Qy   2836 TGTGGGAGGTGCTGATGCTCTGTAGGGCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGG 2895
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Db   2579 TGTGGGAGGTGCTGATGCTCTGTAGGGCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGG 2638

Qy   2896 TCATCGAGAACGCGGGGGAGTTCTTCCGGGACCAGGGCCGGCAGGTGTACCTGTCCCGGC 2955
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Db   2699 CGCCTGCCTGCCCGCAGGGCCTATATGAGCTGATGCTTCGGTGCTGGAGCCGGGAGTCTG 2758

Qy   3016 AGCAGCGACCACCCTTTTCCCAGCTGCATCGGTTCCTGGCAGAGGATGCACTCAACACGG 3075
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2759 AGCAGCGACCACCCTTTTCCCAGCTGCATCGGTTCCTGGCAGAGGATGCACTCAACACGG 2818

Qy   3076 TGTGAATCACACATCCAGCTGCCCCTCCCTCAGGGAGTGATCCAGGGGAAGCCAGTGACA 3135
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Db   2819 TGTGAATCACACATCCAGCTGCCCCTCCCTCAGGGAGTGATCCAGGGGAAGCCAGTGACA 2878

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Db   2879 CTAAAACAAGAGGACACAATGGCACCTCTGCCCTTCCCCTCCCGACAGCCCATCACCTCT 2938

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Db   2939 AATAGAGGCAGTGAGACTGC---------------------------------------- 2958

Qy   3256 CCCTTCCTGGACACACTCTCATGTCCCCTTCCTGTTCTTCCTTCCTAGAAGCCCCTGTCG 3315
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Qy   3316 CCCACCCAGCTGGTCCTGTGGATGGGATCCTCTCCACCCTCCTCTAGCCATCCCTTGGGG 3375
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Qy   3376 AAGGGTGGGGAGAAATATAGGATAGACACTGGACATGGCCCATTGGAGCACCTGGGCCCC 3435
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Db   3033 AAGGGTGGGGAGAAATATAGGATAGACACTGGACATGGCCCATTGGAGCACCTGGGCCCC 3092

Qy   3436 ACTGGACAACACTGATTCCTGGAGAGGTGGCTGCG-CCCCAGCTTCTCTCTCCCTGTCAC 3494
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Db   3093 ACTGGACAACACTGATTCCTGGAGAGGTGGCTGCGCCCCCAGCTTCTCTCTCCCTGTCAC 3152

Qy   3495 ACACTGGACCCCACTGGCTGAGAATCTGGGGGTGAGGAGGACAAGAAGGAGAGGAAAATG 3554
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Db   3153 ACACTGGACCCCACTGGCTGAGAATCTGGGGGTGAGGAGGACAAGAAGGAGAGGAAAATG 3212

Qy   3555 TTTCCTTGTGCCTGCTCCTGTACTTGTCCTCAGCTTGGGCTTCTTCCTCCTCCATCACCT 3614
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Db   3213 TTTCCTTGTGCCTGCTCCTGTACTTGTCCTCAGCTTGGGCTTCTTCCTCCTCCATCACCT 3272

Qy   3615 GAAACACTGGACCTGGGGGTAGCCCCGCCCCAGCCCTCAGTCACCCCCACTTCCCACTTG 3674
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Db   3273 GAAACACTGGACCTGGGGGTAGCCCCGCCCCAGCCCTCAGTCACCCCCACTTCCCACCTG 3332

Qy   3675 CAGTCTTGTAGCTAGAACTTCTCTAAGCCTATACGTTTCTGTGGAGTAAATATTGGGATT 3734
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3333 CAGTCTTGTAGCTAGAACTTCTCTAAGCCTATACGTTTCTGTGGAGTAAATATTGGGATT 3392

Qy   3735 GGGGGGAAAGAGGGAGCAACGGCCCATAGCCTTGGGGTTGGACATCTCTAGTGTAGCTGC 3794
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Db   3393 GGGGGGAAAGAGGGAGCAACGGCCCATAGCCTTGGGGTTGGACATCTCTAGTGTAGCTGC 3452

Qy   3795 CACATTGATTTTTCTATAATCACTTGGGGTTTGTACATTTTTGGGGGGAGAGACACAGAT 3854
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Db   3453 CACATTGATTTTTCTATAATCACTTGGGGTTTGTACATTTTTGGGGGGAGAGACACAGAT 3512

Qy   3855 TTTTACACTAATATATGGACCTAGCTTGAGGCAATTTTAATCCCCTGCACTAGGCAGGTA 3914
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Db   3513 TTTTACACTAATATATGGACCTAGCTTGAGGCAATTTTAATCCCCTGCACTAGGCAGGTA 3572

Qy   3915 ATAATAAAGGTTGAGTTTTCCACAAAAAAAAAAAAAAAA 3953
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Db   3573 ATAATAAAGGTTGAGTTTTCCACAAAAAAAAAAAAAAAA 3611
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RESULT 3
US-08-170-558-3
; Sequence 3, Application US/08170558
; Patent No. 6001621
; GENERAL INFORMATION:
; APPLICANT: Godowski, Paul J.
; APPLICANT: Mark, Melanie R.
; APPLICANT: Scadden, David T.
; APPLICANT: Baker, Kevin P.

```
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/170,558
;      FILING DATE:  20-DEC-1993
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C1
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3637 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-170-558-3
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Query Match 87.1%; Score 3451; DB 3; Length 3637;
Best Local Similarity 97.0%; Pred. No. 0;
Matches 3589; Conservative 0; Mismatches 5; Indels 105; Gaps 3;

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Db   137 ATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAGG 196

Qy   436 ACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCCG 495
Db   197 ACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCCG 256

Qy   496 CCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCGG 555
Db   257 CCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCGG 316

Qy   556 TGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTCCACCTGGTGGCTC 615
Db   317 TGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTGCACCTGGTGGCTC 376

Qy   616 TGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTACC 675
Db   377 TGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTACC 436

Qy   676 GGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAGG 735
Db   437 GGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAGG 496

Qy   736 AGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCCA 795
Db   497 AGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCCA 556

Qy   796 TGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGTGTCTGTCTGC 855
Db   557 TGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGCGTCTGTCTGC 616

Qy   856 GGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTGG 915
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Qy   916 GGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACATA 975
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Db   737 CCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGG 796

Qy  1036 ATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGA 1095
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Qy  1096 GCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGG 1155
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Qy  1156 CCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCTG 1215
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Qy  1216 GCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGC 1275
Db   977 GCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGC 1036

Qy  1276 GCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTG 1335
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Qy  1336 GCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCT 1395
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Qy  1396 TCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCA 1455
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Qy  1456 CCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGG 1515
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Qy  1516 AGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCCATCC 1575
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Qy  1576 TCATCGGCTGCCTGGTGGCCATCATCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCT 1635
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Qy  1636 GGCGGCTGCACTGGCGCAGGCTCCTCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGC 1695
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Qy  1696 TGACGGTTCACCTCTCTGTCCCTGGGGACACTATCCTCATCAACAACCGCCCAGGTCCTA 1755
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Qy  1756 GAGAGCCACCCCCGTACCAGGAGCCCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCT 1815
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Db   1697 ACACCCAGGCCTACAGTGGGGACTATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGC 1756

Qy   1996 CCCCACCTCCCCAGAACAGCGTCCCCCATTATGCCGAGGCTGACATTGTTACCCTGCAGG 2055
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Qy   2116 GGCCCCCCAGAGTGGATTTCCCTCGATCTCGACTCCGCTTCAAGGAGAAGCTTGGCGAGG 2175
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Qy   2176 GCCAGTTTGGGGAGGTGCACCTGTGTGAGGTCGACAGCCCTCAAGATCTGGTCAGTCTTG 2235
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Db   1937 GCCAGTTTGGGGAGGTGCACCTGTGTGAGGTCGACAGCCCTCAAGATCTGGTCAGTCTTG 1996

Qy   2236 ATTTCCCCCTTAATGTGCGTAAGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGC 2295
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Qy   2296 CAGATGCCACCAAGAATGCCAGCTTCTCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGG 2355
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Db   2099 TGAAGATCATGTCGAGGCTCAAGGACCCAAACATCATTCGGCTGCTGGGCGTGTGTGTGC 2158

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Db   2159 AGGACGACCCCCTCTGCATGATTACTGACTACATGGAGAACGGCGACCTCAACCAGTTCC 2218

Qy   2476 TCAGTGCCCACCAGCTGGAGGACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTG 2535
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Db   2219 TCAGTGCCCACCAGCTGGAGGACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTG 2278

Qy   2536 CGCAGGGGCCCACCATCAGCTACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCG 2595
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Db   2279 CGCAGGGGCCCACCATCAGCTACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCG 2338

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Db   2339 GCATGCGCTATCTGGCCACACTCAACTTTGTACATCGGGACCTGGCCACGCGGAACTGCC 2398

Qy   2656 TAGTTGGGGAAAATTTCACCATCAAAATCGCAGACTTTGGCATGAGCCGGAACCTCTATG 2715
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2399 TAGTTGGGGAAAATTTCACCATCAAAATCGCAGACTTTGGCATGAGCCGGAACCTCTATG 2458

Qy   2716 CTGGGGACTATTACCGTGTGCAGGGCCGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGG 2775
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   2459 CTGGGGACTATTACCGTGTGCAGGGCCGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGG 2518

Qy   2776 AGTGCATCCTCATGGGGAAGTTCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCC 2835
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Qy   2836 TGTGGGAGGTGCTGATGCTCTGTAGGGCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGG 2895
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Qy   2896 TCATCGAGAACGCGGGGGGAGTTCTTCCGGGACCAGGGCCGGCAGGTGTACCTGTCCCGGC 2955
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Qy   2956 CGCCTGCCTGCCCGCAGGGCCTATATGAGCTGATGCTTCGGTGCTGGAGCCGGGAGTCTG 3015
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Qy   3495 ACACTGGACCCCACTGGCTGAGAATCTGGGGGTGAGGAGGACAAGAAGGAGAGGAAAATG 3554
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Db   3153 ACACTGGACCCCACTGGCTGAGAATCTGGGGGTGAGGAGGACAAGAAGGAGAGGAAAATG 3212

Qy   3555 TTTCCTTGTGCCTGCTCCTGTACTTGTCCTCAGCTTGGGCTTCTTCCTCCTCCATCACCT 3614
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Db   3273 GAAACACTGGACCTGGGGGTAGCCCCGCCCCAGCCCTCAGTCACCCCCACTTCCCACCTG 3332

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Qy   3795 CACATTGATTTTTCTATAATCACTTGGGGTTTGTACATTTTTGGGGGGAGAGACACAGAT 3854
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RESULT   4
US-08-447-314-3
; Sequence 3, Application US/08447314
; Patent No. 6087144
;  GENERAL INFORMATION:
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/447,314
;      FILING DATE:  22-MAY-1995
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/170558
;      FILING DATE:  20-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C1D2
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3637 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-447-314-3


  Query Match             87.1%;  Score 3451;  DB 3;  Length 3637;
  Best Local Similarity   97.0%;  Pred. No. 0;
  Matches 3589;  Conservative    0;  Mismatches    5;  Indels  105;  Gaps    3;

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Db   1037 GCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTG 1096

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Db   1097 GCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCT 1156

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        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1217 CCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGG 1276

Qy   1516 AGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCCATCC 1575
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1277 AGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGCCCGAGGGGAGCCCGACCGCCATCC 1336

Qy   1576 TCATCGGCTGCCTGGTGGCCATCATCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCT 1635
        ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1337 TCATCGGCTGCCTGGTGGCCATCATCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCT 1396

Qy   1636 GGCGGCTGCACTGGCGCAGGCTCCTCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGC 1695
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Qy  1696 TGACGGTTCACCTCTCTGTCCCTGGGGACACTATCCTCATCAACAACCGCCCAGGTCCTA 1755

Db  1457 TGACGGTTCACCTCTCTGTCCCTGGGGACACTATCCTCATCAACAACCGCCCAGGTCCTA 1516

Qy  1756 GAGAGCCACCCCCGTACCAGGAGCCCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCT 1815

Db  1517 GAGAGCCACCCCCGTACCAGGAGCCCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCT 1576

Qy  1816 GTGTCCCCAATGGCTCTGCGTTGCTGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCA 1875

Db  1577 GTGTCCCCAATGGCTCTGCGTTGCTGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCA 1636

Qy  1876 CTTACGCCCGTCCCCCTCGAGGCCCGGGCCCCCCCACACCCGCCTGGGCCAAACCCACCA 1935

Db  1637 CTTACGCCCGTCCCCCTCGAGGCCCGGGCCCCCCCACACCCGCCTGGGCCAAACCCACCA 1696

Qy  1936 ACACCCAGGCCTACAGTGGGGACTATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGC 1995

Db  1697 ACACCCAGGCCTACAGTGGGGACTATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGC 1756

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Db  1757 CCCCACCTCCCCAGAACAGCGTCCCCCATTATGCCGAGGCTGACATTGTTACCCTGCAGG 1816

Qy  2056 GCGTCACCGGGGGCAACACCTATGCTGTGCCTGCACTGCCCCCAGGGGCAGTCGGGGATG 2115

Db  1817 GCGTCACCGGGGGCAACACCTATGCTGTGCCTGCACTGCCCCCAGGGGCAGTCGGGGATG 1876

Qy  2116 GGCCCCCCAGAGTGGATTTCCCTCGATCTCGACTCCGCTTCAAGGAGAAGCTTGGCGAGG 2175

Db  1877 GGCCCCCCAGAGTGGATTTCCCTCGATCTCGACTCCGCTTCAAGGAGAAGCTTGGCGAGG 1936

Qy  2176 GCCAGTTTGGGGAGGTGCACCTGTGTGAGGTCGACAGCCCTCAAGATCTGGTCAGTCTTG 2235

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Qy  2236 ATTTCCCCCTTAATGTGCGTAAGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGC 2295

Db  1997 ATTTCCCCCTTAATGTGCGTAAGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGC 2056

Qy  2296 CAGATGCCACCAAGAATGCCAGCTTCTCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGG 2355

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Qy  2356 TGAAGATCATGTCGAGGCTCAAGGACCCCAACATCATTCGGCTGCTGGGCGTGTGTGTGC 2415

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Db  2159 AGGACGACCCCCTCTGCATGATTACTGACTACATGGAGAACGGCGACCTCAACCAGTTCC 2218

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Db  2519 AGTGCATCCTCATGGGGAAGTTCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCC 2578

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Qy  3076 TGTGAATCACACATCCAGCTGCCCCTCCCTCAGGGAGTGATCCAGGGGAAGCCAGTGACA 3135

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Qy  3136 CTAAAACAAGAGGACACAATGGCACCTCTGCCCTTCCCCTCCCGACAGCCCATCACCTCT 3195

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Qy  3196 AATAGAGGCAGTGAGACTGCAGGTGGGCTGGGCCCACCCAGGGAGCTGATGCCCCTTCTC 3255

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Db  2959 ----------------------------------------AGAAGCCCCTGTCG 2972

Qy  3316 CCCACCCAGCTGGTCCTGTGGATGGGATCCTCTCCACCCTCCTCTAGCCATCCCTTGGGG 3375

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Qy  3376 AAGGGTGGGGAGAAATATAGGATAGACACTGGACATGGCCCATTGGAGCACCTGGGCCCC 3435

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Db  3093 ACTGGACAACACTGATTCCTGGAGAGGTGGCTGCGCCCCCAGCTTCTCTCTCCCTGTCAC 3152

Qy  3495 ACACTGGACCCCACTGGCTGAGAATCTGGGGGTGAGGAGGACAAGAAGGAGAGGAAAATG 3554

Db  3153 ACACTGGACCCCACTGGCTGAGAATCTGGGGGTGAGGAGGACAAGAAGGAGAGGAAAATG 3212

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Db  3273 GAAACACTGGACCTGGGGGTAGCCCCGCCCCAGCCCTCAGTCACCCCCACTTCCCACCTG 3332

Qy  3675 CAGTCTTGTAGCTAGAACTTCTCTAAGCCTATACGTTTCTGTGGAGTAAATATTGGGATT 3734

Db  3333 CAGTCTTGTAGCTAGAACTTCTCTAAGCCTATACGTTTCTGTGGAGTAAATATTGGGATT 3392

Qy  3735 GGGGGGAAAGAGGGAGCAACGGCCCATAGCCTTGGGGTTGGACATCTCTAGTGTAGCTGC 3794

Db  3393 GGGGGGAAAGAGGGAGCAACGGCCCATAGCCTTGGGGTTGGACATCTCTAGTGTAGCTGC 3452

Qy  3795 CACATTGATTTTTCTATAATCACTTGGGGTTTGTACATTTTTGGGGGGAGAGACACAGAT 3854

Db  3453 CACATTGATTTTTCTATAATCACTTGGGGTTTGTACATTTTTGGGGGGAGAGACACAGAT 3512
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Qy   3855 TTTTACACTAATATATGGACCTAGCTTGAGGCAATTTTAATCCCCTGCACTAGGCAGGTA 3914
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Db   3513 TTTTACACTAATATATGGACCTAGCTTGAGGCAATTTTAATCCCCTGCACTAGGCAGGTA 3572

Qy   3915 ATAATAAAGGTTGAGTTTTCCACAAAAAAAAAAAAAAAA 3953
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Db   3573 ATAATAAAGGTTGAGTTTTCCACAAAAAAAAAAAAAAAA 3611


RESULT   5
US-08-445-461-3
| Sequence 3, Application US/08445461
; Patent No. 6096527
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/445,461
;      FILING DATE:  22-MAY-1995
;      CLASSIFICATION:  530
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/170558
;      FILING DATE:  20-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C3
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3637 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-445-461-3


  Query Match             87.1%;  Score 3451;  DB 3;  Length 3637;
  Best Local Similarity   97.0%;  Pred. No. 0;
  Matches 3589;  Conservative    0;  Mismatches    5;  Indels  105;  Gaps    3;

Qy    256 GTTGGACTTGAAGGAATGCCAAGAGATGCTGCCCCCACCCCCTTAGGCCCGAGGGATCAG 315
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     17 GTTGGACTTGAAGGAATGCCAAGAGATGCTGCCCCCACCCCCTTAGGCCCGAGGGATCAG 76

Qy    316 GAGCTATGGGACCAGAGGCCCTGTCATCTTTACTGCTGCTGCTCTTGGTGGCAAGTGGAG 375
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     77 GAGCTATGGGACCAGAGGCCCTGTCATCTTTACTGCTGCTGCTCTTGGTGGCAAGTGGAG 136

Qy    376 ATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAGG 435
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Db    137 ATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAGG 196

Qy    436 ACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCCG 495
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    197 ACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCCG 256

Qy    496 CCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCGG 555
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Db    257 CCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCGG 316

Qy    556 TGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTCCACCTGGTGGCTC 615
          ||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||
Db    317 TGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTGCACCTGGTGGCTC 376

Qy    616 TGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTACC 675
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    377 TGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTACC 436

Qy    676 GGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAGG 735
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    437 GGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAGG 496

Qy    736 AGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCCA 795
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Db    497 AGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCCA 556

Qy    796 TGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGTGTCTGTCTGC 855
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Db    557 TGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGCGTCTGTCTGC 616

Qy    856 GGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTGG 915
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Db    617 GGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTGG 676

Qy    916 GGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACATA 975
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Db    677 GGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACATA 736

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Db    737 CCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGG 796

Qy   1036 ATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGA 1095
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    797 ATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGA 856

Qy   1096 GCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGG 1155
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    857 GCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGG 916

Qy   1156 CCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCTG 1215
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Db    917 CCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCTG 976

Qy   1216 GCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGC 1275
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Db    977 GCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGC 1036

Qy   1276 GCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTG 1335
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1037 GCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTG 1096

Qy   1336 GCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCT 1395
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1097 GCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCT 1156

Qy   1396 TCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCA 1455
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1157 TCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCA 1216

Qy   1456 CCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGG 1515
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Db  1217 CCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGG 1276

Qy  1516 AGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCCATCC 1575

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Qy  1636 GGCGGCTGCACTGGCGCAGGCTCCTCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGC 1695

Db  1397 GGCGGCTGCACTGGCGCAGGCTCCTCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGC 1456

Qy  1696 TGACGGTTCACCTCTCTGTCCCTGGGGACACTATCCTCATCAACAACCGCCCAGGTCCTA 1755

Db  1457 TGACGGTTCACCTCTCTGTCCCTGGGGACACTATCCTCATCAACAACCGCCCAGGTCCTA 1516

Qy  1756 GAGAGCCACCCCCGTACCAGGAGCCCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCT 1815

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Qy  1816 GTGTCCCCAATGGCTCTGCGTTGCTGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCA 1875

Db  1577 GTGTCCCCAATGGCTCTGCGTTGCTGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCA 1636

Qy  1876 CTTACGCCCGTCCCCCTCGAGGCCCGGGCCCCCCCCACACCCGCCTGGGCCAAACCCACCA 1935

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Qy  1936 ACACCCAGGCCTACAGTGGGGACTATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGC 1995

Db  1697 ACACCCAGGCCTACAGTGGGGACTATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGC 1756

Qy  1996 CCCCACCTCCCCAGAACAGCGTCCCCCATTATGCCGAGGCTGACATTGTTACCCTGCAGG 2055

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Qy  2056 GCGTCACCGGGGGCAACACCTATGCTGTGCCTGCACTGCCCCCAGGGGCAGTCGGGGATG 2115

Db  1817 GCGTCACCGGGGGCAACACCTATGCTGTGCCTGCACTGCCCCCAGGGGCAGTCGGGGATG 1876

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Db  1877 GGCCCCCCAGAGTGGATTTCCCTCGATCTCGACTCCGCTTCAAGGAGAAGCTTGGCGAGG 1936

Qy  2176 GCCAGTTTGGGGAGGTGCACCTGTGTGAGGTCGACAGCCCTCAAGATCTGGTCAGTCTTG 2235

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Qy  2236 ATTTCCCCCTTAATGTGCGTAAGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGC 2295

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Qy  2296 CAGATGCCACCAAGAATGCCAGCTTCTCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGG 2355

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Qy  2596 GCATGCGCTATCTGGCCACACTCAACTTTGTACATCGGGACCTGGCCACGCGGAACTGCC 2655

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Qy  2656 TAGTTGGGGAAAATTTCACCATCAAAATCGCAGACTTTGGCATGAGCCGGAACCTCTATG 2715

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Qy  2776 AGTGCATCCTCATGGGGAAGTTCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCC 2835

Db  2519 AGTGCATCCTCATGGGGAAGTTCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCC 2578

Qy  2836 TGTGGGAGGTGCTGATGCTCTGTAGGGCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGG 2895

Db  2579 TGTGGGAGGTGCTGATGCTCTGTAGGGCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGG 2638

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Db  2879 CTAAAACAAGAGGACACAATGGCACCTCTGCCCTTCCCCTCCCGACAGCCCATCACCTCT 2938

Qy  3196 AATAGAGGCAGTGAGACTGCAGGTGGGCTGGGCCCACCCAGGGAGCTGATGCCCCTTCTC 3255

Db  2939 AATAGAGGCAGTGAGACTGC---------------------------------------- 2958

Qy  3256 CCCTTCCTGGACACACTCTCATGTCCCCTTCCTGTTCTTCCTTCCTAGAAGCCCCTGTCG 3315

Db  2959 ----------------------------------------AGAAGCCCCTGTCG 2972

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Db  3033 AAGGGTGGGGAGAAATATAGGATAGACACTGGACATGGCCCATTGGAGCACCTGGGCCCC 3092

Qy  3436 ACTGGACAACACTGATTCCTGGAGAGGTGGCTGCG-CCCCAGCTTCTCTCTCCCTGTCAC 3494

Db  3093 ACTGGACAACACTGATTCCTGGAGAGGTGGCTGCGCCCCCAGCTTCTCTCTCCCTGTCAC 3152

Qy  3495 ACACTGGACCCCACTGGCTGAGAATCTGGGGGTGAGGAGGACAAGAAGGAGAGGAAAATG 3554

Db  3153 ACACTGGACCCCACTGGCTGAGAATCTGGGGGTGAGGAGGACAAGAAGGAGAGGAAAATG 3212

Qy  3555 TTTCCTTGTGCCTGCTCCTGTACTTGTCCTCAGCTTGGGCTTCTTCCTCCTCCATCACCT 3614

Db  3213 TTTCCTTGTGCCTGCTCCTGTACTTGTCCTCAGCTTGGGCTTCTTCCTCCTCCATCACCT 3272

Qy  3615 GAAACACTGGACCTGGGGGTAGCCCCGCCCCAGCCCTCAGTCACCCCCACTTCCCACTTG 3674

Db  3273 GAAACACTGGACCTGGGGGTAGCCCCGCCCCAGCCCTCAGTCACCCCCACTTCCCACCTG 3332
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Qy   3675 CAGTCTTGTAGCTAGAACTTCTCTAAGCCTATACGTTTCTGTGGAGTAAATATTGGGATT 3734
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3333 CAGTCTTGTAGCTAGAACTTCTCTAAGCCTATACGTTTCTGTGGAGTAAATATTGGGATT 3392

Qy   3735 GGGGGGAAAGAGGGAGCAACGGCCCATAGCCTTGGGGTTGGACATCTCTAGTGTAGCTGC 3794
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3393 GGGGGGAAAGAGGGAGCAACGGCCCATAGCCTTGGGGTTGGACATCTCTAGTGTAGCTGC 3452

Qy   3795 CACATTGATTTTTCTATAATCACTTGGGGTTTGTACATTTTTGGGGGGAGAGACACAGAT 3854
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3453 CACATTGATTTTTCTATAATCACTTGGGGTTTGTACATTTTTGGGGGGAGAGACACAGAT 3512

Qy   3855 TTTTACACTAATATATGGACCTAGCTTGAGGCAATTTTAATCCCCTGCACTAGGCAGGTA 3914
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   3513 TTTTACACTAATATATGGACCTAGCTTGAGGCAATTTTAATCCCCTGCACTAGGCAGGTA 3572

Qy   3915 ATAATAAAGGTTGAGTTTTCCACAAAAAAAAAAAAAAAA 3953
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RESULT   6
US-08-445-640-7
; Sequence 7, Application US/08445640
; Patent No. 5709858
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/445,640
;      FILING DATE:  22-MAY-1995
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/170558
;      FILING DATE:  20-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C2
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  7:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1197 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-445-640-7
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  Query Match          30.1%;  Score 1192.2;  DB 1;  Length 1197;
  Best Local Similarity  99.7%;  Pred. No. 6.4e-273;
  Matches 1194;  Conservative    0;  Mismatches    3;  Indels    0;  Gaps    0;

Qy    375 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 434
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Db      1 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 60

Qy    435 GACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCC 494
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     61 GACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCC 120

Qy    495 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 554
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    121 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 180

Qy    555 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTCCACCTGGTGGCT 614
          |||||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db    181 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTGCACCTGGTGGCT 240

Qy    615 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 674
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    241 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 300

Qy    675 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 734
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    301 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 360

Qy    735 GAGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCC 794
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    361 GAGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCC 420

Qy    795 ATGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGTGTCTGTCTG 854
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Db    421 ATGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGCGTCTGTCTG 480

Qy    855 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 914
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    481 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 540

Qy    915 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 974
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    541 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 600

Qy    975 ACCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTG 1034
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    601 ACCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTG 660

Qy   1035 GATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGG 1094
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    661 GATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGG 720

Qy   1095 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 1154
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    721 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 780

Qy   1155 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 1214
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Db    781 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 840

Qy   1215 GGCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATG 1274
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    841 GGCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATG 900

Qy   1275 CGCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTT 1334
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    901 CGCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTT 960

Qy   1335 GGCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTC 1394
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Db    961 GGCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTC 1020

Qy   1395 TTCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGC 1454
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Db    1021 TTCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGC 1080

Qy    1455 ACCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTG 1514
Db    1081 ACCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTG 1140

Qy    1515 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCC 1571
Db    1141 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGCCCGAGGGGAGCCCGACCGCC 1197


RESULT   7
US-08-170-558-7
; Sequence 7, Application US/08170558
; Patent No. 6001621
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/170,558
;      FILING DATE:  20-DEC-1993
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C1
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  7:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1197 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-170-558-7


  Query Match             30.1%;  Score 1192.2;  DB 3;  Length 1197;
  Best Local Similarity   99.7%;  Pred. No. 6.4e-273;
  Matches 1194;  Conservative    0;  Mismatches    3;  Indels    0;  Gaps    0;

Qy     375 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 434
Db       1 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 60

Qy     435 GACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCC 494
Db      61 GACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCC 120

Qy     495 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 554
Db     121 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 180

Qy     555 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTCCACCTGGTGGCT 614
Db     181 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTGCACCTGGTGGCT 240

Qy     615 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 674
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Qy     675 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 734
Db     301 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 360

Qy     735 GAGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCC 794
Db     361 GAGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCC 420

Qy     795 ATGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGTGTCTGTCTG 854
Db     421 ATGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGCGTCTGTCTG 480

Qy     855 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 914
Db     481 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 540

Qy     915 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 974
Db     541 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 600

Qy     975 ACCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTG 1034
Db     601 ACCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTG 660

Qy    1035 GATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGG 1094
Db     661 GATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGG 720

Qy    1095 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 1154
Db     721 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 780

Qy    1155 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 1214
Db     781 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 840

Qy    1215 GGCGGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATG 1274
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Qy    1395 TTCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGC 1454
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Qy    1455 ACCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTG 1514
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Qy    1515 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCC 1571
Db    1141 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGCCCGAGGGGAGCCCGACCGCC 1197


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US-08-447-314-7
; Sequence 7, Application US/08447314
; Patent No. 6087144
;  GENERAL INFORMATION:
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/447,314
;      FILING DATE:  22-MAY-1995
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/170558
;      FILING DATE:  20-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C1D2
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  7:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1197 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-447-314-7
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Query Match 30.1%; Score 1192.2; DB 3; Length 1197;
Best Local Similarity 99.7%; Pred. No. 6.4e-273;
Matches 1194; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

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Qy    375 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 434
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Db      1 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 60

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Db     61 GACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCC 120

Qy    495 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 554
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Db    121 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 180

Qy    555 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTCCACCTGGTGGCT 614
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Db    181 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTGCACCTGGTGGCT 240

Qy    615 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 674
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    241 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 300

Qy    675 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 734
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Db    301 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 360

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Db    361 GAGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCC 420

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          |||||||||||||||||||||||||||||||||||||||||||||| ||||||||
Db    421 ATGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGCGTCTGTCTG 480

Qy    855 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 914
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    481 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 540

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Db    541 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 600

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Db    601 ACCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTG 660

Qy   1035 GATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGG 1094
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Db    661 GATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGG 720

Qy   1095 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 1154
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    721 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 780

Qy   1155 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 1214
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Db    781 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 840

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Db    901 CGCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTT 960

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Db    961 GGCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTC 1020

Qy   1395 TTCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGC 1454
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Db   1021 TTCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGC 1080

Qy   1455 ACCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTG 1514
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1081 ACCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTG 1140

Qy   1515 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCC 1571
          ||||||||||||||||||||||||||||||||||| ||||||||||||||||||||
Db   1141 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGCCCGAGGGGAGCCCGACCGCC 1197
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RESULT  9
US-08-445-461-7
; Sequence 7, Application US/08445461
; Patent No. 6096527
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Scadden, David T.
;    APPLICANT:  Baker, Kevin P.
;    APPLICANT:  Baron, Will F.
;    TITLE OF INVENTION:  Protein Tyrosine Kinases
;    NUMBER OF SEQUENCES:  35
;    CORRESPONDENCE ADDRESS:
```

```
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/445,461
;      FILING DATE:  22-MAY-1995
;      CLASSIFICATION:  530
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/170558
;      FILING DATE:  20-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Hasak, Janet E.
;      REGISTRATION NUMBER:  28,616
;      REFERENCE/DOCKET NUMBER:  854C3
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1896
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  7:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1197 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
US-08-445-461-7


  Query Match             30.1%;  Score 1192.2;  DB 3;  Length 1197;
  Best Local Similarity   99.7%;  Pred. No. 6.4e-273;
  Matches 1194;  Conservative    0;  Mismatches    3;  Indels    0;  Gaps    0;

Qy     375 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 434
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1 GATGCTGACATGAAGGGACATTTTGATCCTGCCAAGTGCCGCTATGCCCTGGGCATGCAG 60

Qy     435 GACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCC 494
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      61 GACCGGACCATCCCAGACAGTGACATCTCTGCTTCCAGCTCCTGGTCAGATTCCACTGCC 120

Qy     495 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 554
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     121 GCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGGATGGGGCCTGGTGCCCCGCAGGGTCG 180

Qy     555 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTCCACCTGGTGGCT 614
           ||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||
Db     181 GTGTTTCCCAAGGAGGAGGAGTACTTGCAGGTGGATCTACAACGACTGCACCTGGTGGCT 240

Qy     615 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 674
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     241 CTGGTGGGCACCCAGGGACGGCATGCCGGGGGCCTGGGCAAGGAGTTCTCCCGGAGCTAC 300

Qy     675 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 734
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     301 CGGCTGCGTTACTCCCGGGATGGTCGCCGCTGGATGGGCTGGAAGGACCGCTGGGGTCAG 360

Qy     735 GAGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCC 794
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     361 GAGGTGATCTCAGGCAATGAGGACCCTGAGGGAGTGGTGCTGAAGGACCTTGGGCCCCCC 420

Qy     795 ATGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGTGTCTGTCTG 854
           |||||||||||||||||||||||||||||||||||||||||||||| |||||||||
Db     421 ATGGTTGCCCGACTGGTTCGCTTCTACCCCCGGGCTGACCGGGTCATGAGCGTCTGTCTG 480

Qy     855 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 914
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Db     481 CGGGTAGAGCTCTATGGCTGCCTCTGGAGGGATGGACTCCTGTCTTACACCGCCCCTGTG 540

Qy     915 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 974
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Db     541 GGGCAGACAATGTATTTATCTGAGGCCGTGTACCTCAACGACTCCACCTATGACGGACAT 600

Qy     975 ACCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTG 1034
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Db     601 ACCGTGGGCGGACTGCAGTATGGGGGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTG 660

Qy    1035 GATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGG 1094
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     661 GATGACTTTAGGAAGAGTCAGGAGCTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGG 720

Qy    1095 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 1154
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Db     721 AGCAACCACAGCTTCTCCAGTGGCTATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGG 780

Qy    1155 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 1214
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     781 GCCTTCCAGGCTATGCAGGTCCACTGTAACAACATGCACACGCTGGGAGCCCGTCTGCCT 840

Qy    1215 GGCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATG 1274
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     841 GGCGGGGTGGAATGTCGCTTCCGGCGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATG 900

Qy    1275 CGCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTT 1334
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     901 CGCCACAACCTAGGGGGCAACCTGGGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTT 960

Qy    1335 GGCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTC 1394
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     961 GGCGGCCGTGTGGCTCGCTTTCTGCAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTC 1020

Qy    1395 TTCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGC 1454
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1021 TTCAGCGAAATCTCCTTCATCTCTGATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGC 1080

Qy    1455 ACCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTG 1514
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1081 ACCTTCCCGCCAGCCCCCTGGTGGCCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTG 1140

Qy    1515 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCC 1571
           |||||||||||||||||||||||||||||||||||| ||||||||||||||||||||
Db    1141 GAGCTGGAGCCCAGAGGCCAGCAGCCCGTGGCCAAGCCCGAGGGGAGCCCGACCGCC 1197


RESULT  10
US-08-336-343A-3
; Sequence 3, Application US/08336343A
; Patent No. 5677144
;  GENERAL INFORMATION:
;    APPLICANT:  Ullrich, Axel
;    APPLICANT:  Alves, Frauke
;    TITLE OF INVENTION:  CCK-2, A No. 5677144el Receptor Tyrosine Kinase
;    NUMBER OF SEQUENCES:  43
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Pennie & Edmonds
;      STREET:  1155 Avenue of the Americas
;      CITY:  New York
;      STATE:  New York
;      COUNTRY:  U.S.A.
;      ZIP:  10036-2711
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/336,343A
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;      FILING DATE:  08-NOV-1994
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Coruzzi, Laura A.
;      REGISTRATION NUMBER:  30,742
;      REFERENCE/DOCKET NUMBER:  7683-065
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (212) 790-9090
;      TELEFAX:  (212) 869-9741/8864
;      TELEX:  66141 PENNIE
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3157 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  both
;      TOPOLOGY:  unknown
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  370..2934
US-08-336-343A-3

  Query Match             16.2%;  Score 642;  DB 1;  Length 3157;
  Best Local Similarity   56.5%;  Pred. No. 1.4e-142;
  Matches 1534;  Conservative    0;  Mismatches  975;  Indels  207;  Gaps   10;

Qy    349 TGCTGCTGCTCTTGGTGGCAAGTGGAGATGCTGACATGAAGGGACATTTTGATCCTGCCA 408
Db    395 TGGTGCTGTTCCTGCTGCTGCCTATCTTGAGTTCTGCAAAAGCTCAGGTTAATCCAGCTA 454

Qy    409 AGTGCCGCTATGCCCTGGGCATGCAGGACCGGACCATCCCAGACAGTGACATCTCTGCTT 468
Db    455 TATGCCGCTATCCTCTGGGCATGTCAGGAGGCCAGATTCCAGATGAGGACATCACAGCTT 514

Qy    469 CCAGCTCCTGGTCAGATTCCACTGCCGCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGG 528
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Qy    529 ATGGGGCCTGGTGCCCCGCAGGGTCGGTGTTTCCCAAGGA---GGAGGAGTACTTGCAGG 585
Db    575 ATGGAGCCTGGTGCCCTGAGATTCCAGTGGAACCTGATGACCTGAAGGAGTTTCTGCAGA 634

Qy    586 TGGATCTACAACGACTCCACCTGGTGGCTCTGGTGGGCACCCAGGGACGGCATGCCGGGG 645
Db    635 TTGACTTGCACACCCTCCATTTTATCACTCTGGTGGGGACCCAGGGGCGCCATGCAGGAG 694

Qy    646 GCCTGGGCAAGGAGTTCTCCCGGAGCTACCGGCTGCGTTACTCCCGGGATGGTCGCCGCT 705
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Qy    706 GGATGGGCTGGAAGGACCGCTGGGGTCAGGAGGTGATCTCAGGCAATGAGGACCCTGAGG 765
Db    755 GGATCTCTTGGCGGAACCGTCATGGGAAACAGGTGCTGGATGGAAATAGTAACCCCTATG 814

Qy    766 GAGTGGTGCTGAAGGACCTTGGGCCCCCCATGGTTGCCCGACTGGTTCGCTTCTACCCCC 825
Db    815 ACATTTTCCTAAAGGACTTGGAGCCGCCCATTGTAGCCAGATTTGTCCGGTTCATTCCAG 874

Qy    826 GGGCTGACCGGGTCATGAGTGTCTGTCTGCGGGTAGAGCTCTATGGCTGCCTCTGGAGGG 885
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Qy    886 ATGGACTCCTGTCTTACACCGCCCCTGTGGGGCAGACAATGTATTTATCTGAGG------ 939
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Qy   1000 GTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGGATGACTTTAGGAAGAGTCAGGAGC 1059
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Qy   1060 TGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGAGCAACCACAGCTTCTCCAGTGGCT 1119
Db   1112 ACCACGTGTGGCCCGGCTATGACTATGTGGGCTGGCGGAACGAGAGTGCCACCAATGGCT 1171

Qy   1120 ATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGGCCTTCCAGGCTATGCAGGTCCACT 1179
Db   1172 ACATTGAGATCATGTTTGAATTTGACCGCATCAGGAATTTCACTACCATGAAGGTCCACT 1231

Qy   1180 GTAACAACATGCACACGCTGGGAGCCCGTCTGCCTGGCGGGGTGGAATGTCGCTTCCGGC 1239
Db   1232 GCAACAACATGTTTGCTAAAGGTGTGAAGATCTTTAAGGAGGTACAGTGCTACTTCCGCT 1291

Qy   1240 GTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGCGCCACAACCTAGGGGGCAACCTGG 1299
Db   1292 CTG---AAGCCAGTGAGTGGGAACCTAATGCCATTTCCTTCCCCCTTGTCCTGGATGACG 1348

Qy   1300 GGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTGGCGGCCGTGTGGCTCGCTTTCTGC 1359
Db   1349 TCAACCCCAGTGCTCGGTTTGTCACGGTGCCTCTCCACCACCGAATGGCCAGTGCCATCA 1408

Qy   1360 AGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCTTCAGCGAAATCTCCTTCATCTCTG 1419
Db   1409 AGTGTCAATACCATTTTGCAGATACCTGGATGATGTTCAGTGAGATCACCTTCCAATCAG 1468

Qy   1420 ATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCACCTTCCCGCCAGCCCCCTGGTGGC 1479
Db   1469 ATGCTGCAATGTACAACAACTCTGAAGCCCTGCCCACCTCTCCTA---------------- 1513

Qy   1480 CGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGGAGCTGGAGCCCAGAGGCCAGCAGC 1539
Db   1514 ----------------------------------------TGGCACCCACAACCTATGATC 1534

Qy   1540 CCGTGGCCAAGGCCGAGGGGAGCCCGACCGCCATCCTCATCGGCTGCCTGGTGGCCATCA 1599
Db   1535 CAATGCTTAAAGTTGATGACAGCAACACTCGGATCCTGATTGGCTGCTTGGTGGCCATCA 1594

Qy   1600 TCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCTGGCGGCTGCACTGGCGCAGGCTCC 1659
Db   1595 TCTTTATCCTCCTGGCCATCATTGTCATCATCCTCTGGAGGCAGTTCTGGCAGAAAATGC 1654

Qy   1660 TCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGCTGACGGTTCACCTCTCTGTCCCTG 1719
Db   1655 TGGAGAAGGCTTCTCGGAGGATGCTGGATGATGAAATGACAGTCAGCCTTTCCCTGCCAA 1714

Qy   1720 GGGACACTATCCTCATCAACAACCGCCCAGGTCCTAGAGAGCCACCCCCGTACCAGGAGC 1779
Db   1715 GTGATTCTAGCATGTTCAACAAT----------------------------AACCGCTCCT 1747

Qy   1780 CCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCTGTGTCCCCAATGGCTCTGCGTTGC 1839
Db   1748 CATCACCTAGTGAACAAGGGTCCAACTCGACTTACGATCGCATCTTT------------- 1794

Qy   1840 TGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCACTTACGCCCGTCCCCCTCGAGGCC 1899
Db   1795 ----------------------------------------------------CCCCTTCGCC 1804

Qy   1900 CGGGCCCCCCCACACCCGCCTGGGCCAAACCCACCAACACCCAGGCCTACAGTGGGGACT 1959
Db   1805 CTGACTACCAGGAGCCATCCAGGCTGATACGAAAACTCCCAGAATTTGCTCCAGGGGAGG 1864

Qy   1960 ATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGCCCCCACCTCCCCAGAACAGCGTCC 2019
Db   1865 AGGAGTCAGGCTGCAGCGGTGTTGTGAAGCCAGTCCAGCCCAGTGGCCCTGAGGGGGTGC 1924

Qy   2020 CCCATTATGCCGAGGCTGACATTGTTACCCTGCAGGGCGTCACCGGGGGCAACACCTATG 2079
Db   1925 CCCACTATGCAGAGGCTGACATAGTGAACCTCCAAGGAGTGACAGGAGGCAACACATACT 1984

Qy   2080 CTGTGCCTGCACTGCCCCCAGGGGCAGTC---GGGGATGGGCCCCCCAGAGTGGATTTCC 2136
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Db    1985 CAGTGCCTGCCGTCACCATGGACCTGCTCTCAGGAAAAGATGTGGCTGTGGAGGAGTTCC 2044

Qy    2137 CTCGATCTCGACTCCGCTTCAAGGAGAAGCTTGGCGAGGGCCAGTTTGGGGAGGTGCACC 2196

Db    2045 CCAGGAAACTCCTAACTTTCAAAGAGAAGCTGGGAGAAGGACAGTTTGGGGAGGTTCATC 2104

Qy    2197 TGTGTGAGGTCGACAGCCCTCAAGATCTGGTCAGTCTTGATTTCCCCCTTAATGTGCGTA 2256

Db    2105 TCTGTGAAGTGGAGGGAATGGAAAAATTCAAAGACAAAGATTTTGCCCTAGATGTCAGTG 2164

Qy    2257 AGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGCCAGATGCCACCAAGAATGCCA 2316

Db    2165 CCAACCAGCCTGTCCTGGTGGCTGTGAAAATGCTCCGAGCAGATGCCAACAAGAATG--- 2221

Qy    2317 GCTTCTCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGGTGAAGATCATGTCGAGGCTCA 2376

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Qy    2377 AGGACCCCAACATCATTCGGCTGCTGGGCGTGTGTGTGCAGGACGACCCCCTCTGCATGA 2436

Db    2267 AGGACCCAAACATCATCCATCTATTATCTGTGTGTATCACTGATGACCCTCTCTGTATGA 2326

Qy    2437 TTACTGACTACATGGAGAACGGCGACCTCAACCAGTTCCTCAGTGCCCACCAGCTGGAGG 2496

Db    2327 TCACTGAATACATGGAGAATGGAGATCTCAATCAGTTTCTTTCCCGCCACGAGCCCCCTA 2386

Qy    2497 ACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTGCGCAGGGGCCCACCATCAGCT 2556

Db    2387 ATTCTTCCTCCAGCG--------------------------ATGTACGCACTGTCAGTT 2419

Qy    2557 ACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCGGCATGCGCTATCTGGCCACAC 2616

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Qy    2617 TCAACTTTGTACATCGGGACCTGGCCACGCGGAACTGCCTAGTTGGGGAAAATTTCACCA 2676

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Qy    2737 AGGGCCGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGGAGTGCATCCTCATGGGGAAGT 2796

Db    2600 AGGGCCGGGCAGTGCTCCCTATCCGCTGGATGTCTTGGGAGAGTATCTTGCTGGGCAAGT 2659

Qy    2797 TCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCCTGTGGGAGGTGCTGATGCTCT 2856

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Qy    2917 TCTTCCGGGACCAGGGCCGGCAGGTGTACCTGTCCCGGCCGCCTGCCTGCCCGCAGGGCC 2976

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Qy    3037 AGCTGCATCGGTTCCT 3052

Db    2900 AAATCCACCTTCTGCT 2915


RESULT  11
US-08-336-343A-5/c
; Sequence 5, Application US/08336343A
; Patent No. 5677144
;  GENERAL INFORMATION:
;    APPLICANT:  Ullrich, Axel
;    APPLICANT:  Alves, Frauke
;    TITLE OF INVENTION:  CCK-2, A No. 5677144el Receptor Tyrosine Kinase
;    NUMBER OF SEQUENCES:  43
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Pennie & Edmonds
;      STREET:  1155 Avenue of the Americas
;      CITY:  New York
;      STATE:  New York
;      COUNTRY:  U.S.A.
;      ZIP:  10036-2711
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/336,343A
;      FILING DATE:  08-NOV-1994
;      CLASSIFICATION:  435
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Coruzzi, Laura A.
;      REGISTRATION NUMBER:  30,742
;      REFERENCE/DOCKET NUMBER:  7683-065
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (212) 790-9090
;      TELEFAX:  (212) 869-9741/8864
;      TELEX:  66141 PENNIE
;  INFORMATION FOR SEQ ID NO:  5:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3157 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  unknown
;    MOLECULE TYPE:  cDNA
;    HYPOTHETICAL:  NO
;    ANTI-SENSE:  NO
US-08-336-343A-5


  Query Match           16.2%;  Score 642;  DB 1;  Length 3157;
  Best Local Similarity  56.5%;  Pred. No. 1.4e-142;
  Matches 1534;  Conservative    0;  Mismatches  975;  Indels  207;  Gaps   10;

Qy     349 TGCTGCTGCTCTTGGTGGCAAGTGGAGATGCTGACATGAAGGGACATTTTGATCCTGCCA 408

Db    2763 TGGTGCTGTTCCTGCTGCTGCCTATCTTGAGTTCTGCAAAAGCTCAGGTTAATCCAGCTA 2704

Qy     409 AGTGCCGCTATGCCCTGGGCATGCAGGACCGGACCATCCCAGACAGTGACATCTCTGCTT 468

Db    2703 TATGCCGCTATCCTCTGGGCATGTCAGGAGGCCAGATTCCAGATGAGGACATCACAGCTT 2644

Qy     469 CCAGCTCCTGGTCAGATTCCACTGCCGCCCGCCACAGCAGGTTGGAGAGCAGTGACGGGG 528

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Qy     529 ATGGGGCCTGGTGCCCCGCAGGGTCGGTGTTTCCCAAGGA---GGAGGAGTACTTGCAGG 585

Db    2583 ATGGAGCCTGGTGCCCTGAGATTCCAGTGGAACCTGATGACCTGAAGGAGTTTCTGCAGA 2524

Qy     586 TGGATCTACAACGACTCCACCTGGTGGCTCTGGTGGGCACCCAGGGACGGCATGCCGGGG 645

Db    2523 TTGACTTGCACACCCTCCATTTTATCACTCTGGTGGGGACCCAGGGGCGCCATGCAGGAG 2464

Qy     646 GCCTGGGCAAGGAGTTCTCCCGGAGCTACCGGCTGCGTTACTCCCGGGATGGTCGCCGCT 705

Db    2463 GTCATGGCATCGAGTTTGCCCCCATGTACAAGATCAATTACAGTCGGGATGGCACTCGCT 2404

Qy     706 GGATGGGCTGGAAGGACCGCTGGGGTCAGGAGGTGATCTCAGGCAATGAGGACCCTGAGG 765

Db    2403 GGATCTCTTGGCGGAACCGTCATGGGAAACAGGTGCTGGATGGAAATAGTAACCCCTATG 2344

Qy     766 GAGTGGTGCTGAAGGACCTTGGGCCCCCCATGGTTGCCCGACTGGTTCGCTTCTACCCCC 825
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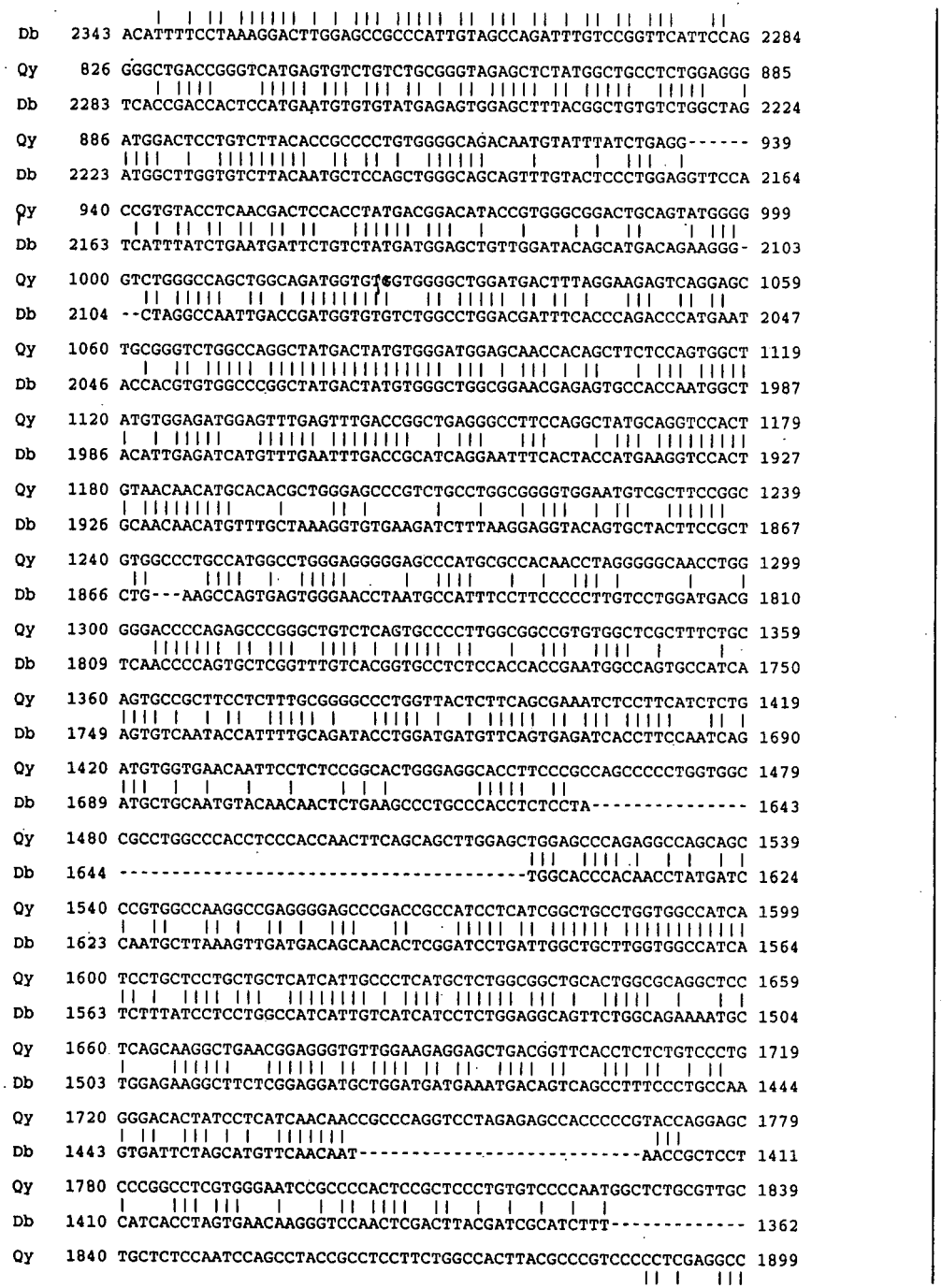
Db 2343 ACATTTTCCTAAAGGACTTGGAGCCGCCCATTGTAGCCAGATTTGTCCGGTTCATTCCAG 2284
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Db 2283 TCACCGACCACTCCATGAATGTGTGTATGAGAGTGGAGCTTTACGGCTGTGTCTGGCTAG 2224
Qy 886 ATGGACTCCTGTCTTACACCGCCCCTGTGGGGCAGACAATGTATTTATCTGAGG------ 939
Db 2223 ATGGCTTGGTGTCTTACAATGCTCCAGCTGGGCAGCAGTTTGTACTCCCTGGAGGTTCCA 2164
Qy 940 CCGTGTACCTCAACGACTCCACCTATGACGGACATACCGTGGGCGGACTGCAGTATGGGG 999
Db 2163 TCATTTATCTGAATGATTCTGTCTATGATGGAGCTGTTGGATACAGCATGACAGAAGGG- 2103
Qy 1000 GTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGGATGACTTTAGGAAGAGTCAGGAGC 1059
Db 2104 --CTAGGCCAATTGACCGATGGTGTGTCTGGCCTGGACGATTTCACCCAGACCCATGAAT 2047
Qy 1060 TGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGAGCAACCACAGCTTCTCCAGTGGCT 1119
Db 2046 ACCACGTGTGGCCCGGCTATGACTATGTGGGCTGGCGGAACGAGAGTGCCACCAATGGCT 1987
Qy 1120 ATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGGCCTTCCAGGCTATGCAGGTCCACT 1179
Db 1986 ACATTGAGATCATGTTTGAATTTGACCGCATCAGGAATTTCACTACCATGAAGGTCCACT 1927
Qy 1180 GTAACAACATGCACACGCTGGGAGCCCGTCTGCCTGGCGGGGTGGAATGTCGCTTCCGGC 1239
Db 1926 GCAACAACATGTTTGCTAAAGGTGTGAAGATCTTTAAGGAGGTACAGTGCTACTTCCGCT 1867
Qy 1240 GTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGCGCCACAACCTAGGGGGCAACCTGG 1299
Db 1866 CTG---AAGCCAGTGAGTGGGAACCTAATGCCATTTCCTTCCCCCTTGTCCTGGATGACG 1810
Qy 1300 GGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTGGCGGCCGTGTGGCTCGCTTTCTGC 1359
Db 1809 TCAACCCCAGTGCTCGGTTTGTCACGGTGCCTCTCCACCACCGAATGGCCAGTGCCATCA 1750
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Db 1644 --TGGCACCCACAACCTATGATC 1624
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Db 1623 CAATGCTTAAAGTTGATGACAGCAACACTCGGATCCTGATTGGCTGCTTGGTGGCCATCA 1564
Qy 1600 TCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCTGGCGGCTGCACTGGCGCAGGCTCC 1659
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Qy 1660 TCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGCTGACGGTTCACCTCTCTGTCCCTG 1719
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Qy 1780 CCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCTGTGTCCCCAATGGCTCTGCGTTGC 1839
Db 1410 CATCACCTAGTGAACAAGGGTCCAACTCGACTTACGATCGCATCTTT------------ 1362
Qy 1840 TGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCACTTACGCCCGTCCCCCTCGAGGCC 1899
Db 1363 --CCCCTTCGCC 1354
Qy 1900 CGGGCCCCCCCACACCCGCCTGGGCCAAACCCACCAACACCCAGGCCTACAGTGGGGACT 1959
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Qy 1960 ATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGCCCCCACCTCCCCAGAACAGCGTCC 2019
Db 1293 AGGAGTCAGGCTGCAGCGGTGTTGTGAAGCCAGTCCAGCCCAGTGGCCCTGAGGGGGTGC 1234
Qy 2020 CCCATTATGCCGAGGCTGACATTGTTACCCTGCAGGGCGTCACCGGGGGCAACACCTATG 2079
Db 1233 CCCACTATGCAGAGGCTGACATAGTGAACCTCCAAGGAGTGACAGGAGGCAACACATACT 1174
Qy 2080 CTGTGCCTGCACTGCCCCCAGGGGCAGTC---GGGGATGGGCCCCCCAGAGTGGATTTCC 2136
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Qy 2137 CTCGATCTCGACTCCGCTTCAAGGAGAAGCTTGGCGAGGGCCAGTTTGGGGAGGTGCACC 2196
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Db 1053 TCTGTGAAGTGGAGGGAATGGAAAAATTCAAAGACAAAGATTTTGCCCTAGATGTCAGTG 994
Qy 2257 AGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGCCAGATGCCACCAAGAATGCCA 2316
Db 993 CCAACCAGCCTGTCCTGGTGGCTGTGAAAATGCTCCGAGCAGATGCCAACAAGAATG--- 935
Qy 2317 GCTTCTCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGGTGAAGATCATGTCGAGGCTCA 2376
Db 936 --------------CCAGGAATGATTTTCTTAAGGAGATAAAGATCATGTCTCGGCTCA 892
Qy 2377 AGGACCCCAACATCATTCGGCTGCTGGGCGTGTGTGTGCAGGACGACCCCCTCTGCATGA 2436
Db 891 AGGACCCAAACATCATCCATCTATTATCTGTGTGTATCACTGATGACCCTCTCTGTATGA 832
Qy 2437 TTACTGACTACATGGAGAACGGCGACCTCAACCAGTTCCTCAGTGCCCACCAGCTGGAGG 2496
Db 831 TCACTGAATACATGGAGAATGGAGATCTCAATCAGTTTCTTTCCCGCCACGAGCCCCCTA 772
Qy 2497 ACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTGCGCAGGGGCCCACCATCAGCT 2556
Db 771 ATTCTTCCTCCAGCG--------------------------ATGTACGCACTGTCAGTT 739
Qy 2557 ACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCGGCATGCGCTATCTGGCCACAC 2616
Db 738 ACACCAATCTGAAGTTTATGGCTACCCAAATTGCCTCTGGCATGAAGTACCTTTCCTCTC 679
Qy 2617 TCAACTTTGTACATCGGGACCTGGCCACGCGGAACTGCCTAGTTGGGGAAAATTTCACCA 2676
Db 678 TTAATTTTGTTCACCGAGATCTGGCCACACGAAACTGTTTAGTGGGTAAGAACTACACAA 619
Qy 2677 TCAAAATCGCAGACTTTGGCATGAGCCGGAACCTCTATGCTGGGGACTATTACCGTGTGC 2736
Db 618 TCAAGATAGCTGACTTTGGAATGAGCAGGAACCTGTACAGTGGTGACTATTACCGGATCC 559
Qy 2737 AGGGCCGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGGAGTGCATCCTCATGGGGAAGT 2796
Db 558 AGGGCCGGGCAGTGCTCCCTATCCGCTGGATGTCTTGGGAGAGTATCTTGCTGGGCAAGT 499
Qy 2797 TCACGACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCCTGTGGGAGGTGCTGATGCTCT 2856
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Qy 2857 GTAGGGCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGGTCATCGAGAACGCGGGGGAGT 2916
Db 438 GTCAAGAACAGCCCTATTCCCAGCTGTCAGATGAACAGGTTATTGAGAATACTGGAGAGT 379
Qy 2917 TCTTCCGGGACCAGGGCCGGCAGGTGTACCTGTCCCGGCCGCCTGCCTGCCCGCAGGGCC 2976
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Qy   2977 TATATGAGCTGATGCTTCGGTGCTGGAGCCGGGAGTCTGAGCAGCGACCACCCTTTTCCC 3036
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Db    318 TGTATAAGCTGATGCTCAGCTGCTGGAGAAGAGATACGAAGAACCGTCCCTCATTCCAAG 259

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RESULT 12
US-08-456-647B-19
; Sequence 19, Application US/08456647B
; Patent No. 5811516
; GENERAL INFORMATION:
;   APPLICANT: Lemke Ph.D. et al, Greg E.
;   TITLE OF INVENTION: PROTEIN-TYROSINE KINASE GENES
;   NUMBER OF SEQUENCES: 54
;   CORRESPONDENCE ADDRESS:
;     ADDRESSEE: Fish & Richardson P.C.
;     STREET: 4225 Executive Square, Suite 1400
;     CITY: La Jolla
;     STATE: CA
;     COUNTRY: US
;     ZIP: 92037
;   COMPUTER READABLE FORM:
;     MEDIUM TYPE: Floppy disk
;     COMPUTER: IBM PC compatible
;     OPERATING SYSTEM: PC-DOS/MS-DOS
;     SOFTWARE: PatentIn Release #1.0, Version #1.25
;   CURRENT APPLICATION DATA:
;     APPLICATION NUMBER: US/08/456,647B
;     FILING DATE: 02-JUN-1995
;     CLASSIFICATION: 530
;   PRIOR APPLICATION DATA:
;     APPLICATION NUMBER: US 08/237,401
;     FILING DATE: 02-MAY-1994
;   PRIOR APPLICATION DATA:
;     APPLICATION NUMBER: US 07/884,486
;     FILING DATE: 15-MAY-1992
;   ATTORNEY/AGENT INFORMATION:
;     NAME: Wetherell Ph.D., John R.
;     REGISTRATION NUMBER: 31,678
;     REFERENCE/DOCKET NUMBER: 07251/007002
;   TELECOMMUNICATION INFORMATION:
;     TELEPHONE: (619) 678-5070
;     TELEFAX: (619) 678-5099
;  INFORMATION FOR SEQ ID NO: 19:
;   SEQUENCE CHARACTERISTICS:
;     LENGTH: 3120 base pairs
;     TYPE: nucleic acid
;     STRANDEDNESS: single
;     TOPOLOGY: linear
;   MOLECULE TYPE: DNA
;   IMMEDIATE SOURCE:
;     CLONE: Tyro-10
;   FEATURE:
;     NAME/KEY: CDS
;     LOCATION: 485..3047
US-08-456-647B-19


  Query Match             16.1%;  Score 639.8;  DB 1;  Length 3120;
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  Matches 1552;  Conservative    0;  Mismatches  987;  Indels  210;  Gaps   11;

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Db    569 ATATGCCGCTATCCTCTGGGCATGTCAGGAGGCCACATTCCAGATGAGGACATCACAGCC 628

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Db    629 TCAAGTCAGTGGTCAGAATCCACGGCTGCCAAATATGGGAGGCTGGACTCTGAAGAAGGA 688

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Db    689 GATGGAGCCTGGTGTCCTGAGATTCCAGTGCAACCCGATGACCTGAAGGAATTTCTGCAG 748

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Db    749 ATTGACTTGCGAACCCTACACTTTATCACTCTTGTGGGGACCCAGGGGCGCCATGCAGGG 808

Qy    645 GGCCTGGGCAAGGAGTTCTCCCGGAGCTACCGGCTGCGTTACTCCCGGGATGGTCGCCGC 704
          || |  |||| || ||  | |  |  ||| | |    |||  |||||||  | |||
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Db   1226 TACCACGTGTGGCCTGGCTATGACTACGTGGGATGGCGGAACGAAAGTGCTACCAACGGT 1285

Qy   1119 TATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGGCCTTCCAGGCTATGCAGGTCCAC 1178
          |   | |||||   |||||| ||||||||  | |||   ||     | ||| ||||||||
Db   1286 TTCATTGAGATCATGTTTGAATTTGACCGAATCAGGAATTTTACTACCATGAAGGTCCAC 1345

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          || |||||||||     |    || |      |   |   | |||  | ||   ||| ||
Db   1346 TGCAACAACATGTTTGCTAAAGGTGTGAAGATTTTTAAGGAGGTCCAGTGCTACTTTCGC 1405

Qy   1239 CGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGCGCCACAACCTAGGGGGCAACCTG 1298
             |    |||| |  |||||         |  |   |     ||| |       ||
Db   1406 TCGG---AAGCCAGCGAGTGGGAACCCACTGCTGTCTACTTTCCCCTGGTCCTGGACGAT 1462

Qy   1299 GGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTGGCGGCCGTGTGGCTCGCTTTCTG 1358
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Db   1463 GTGAACCCCAGTGCCCGGTTTGTCACGGTGCCCCTCCACCACCGAATGGCCAGTGCCATC 1522

Qy   1359 CAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCTTCAGCGAAATCTCCTTCATCTCT 1418
           ||||||  | ||  ||||| |   | ||| |  | |||||||| ||| | |||   ||
Db   1523 AAGTGCCAATACCATTTTGCCGACACGTGGATGATGTTCAGCGAGATCACTTTCCAATCA 1582

Qy   1419 GATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCACCTTCCCGCCAGCCCCCTGGTGG 1478
          ||||  |  |  || |   | | |        |||||  ||
Db   1583 GATGCTGCAATGTATAACAACTCTGGAGCCCTTCCCACCTCTCCTA-------------- 1628

Qy   1479 CCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGGAGCTGGAGCCCAGAGGCCAGCAG 1538
                                                  |||  ||||    | |  |
Db   1629 ----------------------------------------TGGCACCCACCACCTATGAT 1648
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Qy   1539 CCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCCATCCTCATCGGCTGCCTGGTGGCCATC 1598
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Db   1649 CCCATGCTTAAAGTTGATGATAGCAACACTCGGATCCTGATTGGTTGCTTGGTGGCCATC 1708

Qy   1599 ATCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCTGGCGGCTGCACTGGCGCAGGCTC 1658
          ||| |  |||||||  ||||| | | |||| || ||| ||| |  |||||  | | |
Db   1709 ATCTTCATCCTGCTGGCTATCATCGTCATCATCCTGTGGAGGCAGTTCTGGCAGAAGATG 1768

Qy   1659 CTCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGCTGACGGTTCACCTCTCTGTCCCT 1718
          ||    ||||||  |||||| || |||| || ||  |||| ||   ||| ||  | ||
Db   1769 CTAGAAAAGGCTTCACGGAGGATGCTGGATGATGAAATGACAGTCAGCCTTTCCCTGCCC 1828

Qy   1719 GGGGACACTATCCTCATCAACAACCGCCCAGGTCCTAGAGAGCCACCCCCGTACCAGGAG 1778
           | ||  | | |  |||| ||| ||       |        ||||     | ||||||
Db   1829 AGCGAGTCCAGCATGTTCAATAACAACCGCTCCTC------ATCACCAAGTGAACAGGAG 1882

Qy   1779 CCCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCTGTGTCCCCAATGGCTCTGCGTTG 1838
           ||    ||  |   |||   | | ||  || |  || |  | | |   |  |
Db   1883 TCCAACTCTACTTATGATCGAATCTTCCCCCTTCGCCCTGACTACCAGGAGCCATCCAGA 1942

Qy   1839 CTGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCACTTACGCCCGTCCCCCTCGAGGC 1898
          |||  ||
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Qy   1899 CCGGGCCCCCCCACACCCGCCTGGGCCAAACCCACCAACACCCAGGCCTACAGTGGGGAC 1958
                                  | ||| |  |||        ||   || || |
Db   1949 -------------------------CGAAAGCTTCCAGAGTTTGCTCCAGGAGAGGAGGA 1983

Qy   1959 TATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGCCCCCACCTCCCCAGAACAGCGTC 2018
           | || ||    |     |  |||   |     |  ||   |   |   | ||||
Db   1984 GTCAGGGTGCAGTGGTGTTGTGAAGCCGGCC-----CAGCCCAATGGACCTGAGGGCGTG 2038

Qy   2019 CCCCATTATGCCGAGGCTGACATTGTTACCCTGCAGGGCGTCACCGGGGGCAACACCTAT 2078
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Qy   2079 GCTGTGCCTGCACTGCCCCCAG---GGGCAGTCGGGGATGGGCCCCCCAGAGTGGATTTC 2135
           |||||||||  |  ||   |    |  |  ||| | |      |   || |||
Db   2099 TGTGTGCCTGCTGTAACCATGGATCTGCTATCGGGGAAAGATGTGGCTGTGGAAGAGTTC 2158

Qy   2136 CCTCGATCTCGACTCCGCTTCAAGGAGAAGCTTGGCGAGGGCCAGTTTGGGGAGGTGCAC 2195
          ||  |    |   |||||||||||||||| || || ||||||||||||||||| ||
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Qy   2196 CTGTGTGAGGTCGACAGCCCTCAAGATCTGGTCAGTCTTGATTTCCCCCTTAATGTGCGT 2255
          || ||||| || ||  |     || |  |           |||||  | ||  ||||  ||
Db   2219 CTCTGTGAAGTGGAGGGAATGGAAAAATTCAAAGACAAAGATTTTGCACTAGATGTCAGT 2278

Qy   2256 AAGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGCCAGATGCCACCAAGAATGCC 2315
               || ||| | ||||| || || || ||  | ||  |||||||| ||||||||
Db   2279 GCCAACCAGCCTGTCCTGGTGGCCGTGAAAATGCTCCGAGCAGATGCCAACAAGAATG-- 2336

Qy   2316 AGCTTCTCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGGTGAAGATCATGTCGAGGCTC 2375
                     |||||||||||||| || || ||| | ||||||||||  |||||
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Qy   2376 AAGGACCCCAACATCATTCGGCTGCTGGGCGTGTGTGTGCAGGACGACCCCCTCTGCATG 2435
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Db   2381 AAGGACCCAAACATCATCCGTCTCTTAGCTGTGTGCATCACTGAGGACCCGCTCTGCATG 2440

Qy   2436 ATTACTGACTACATGGAGAACGGCGACCTCAACCAGTTCCTCAGTGCCCACCAGCTGGAG 2495
          || || ||  |||||||||| || || || || ||||| ||    |  |||| |||    |
Db   2441 ATCACGGAATACATGGAGAATGGAGATCTTAATCAGTTTCTTTCTCGCCACGAGCCTCTG 2500

Qy   2496 GACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTGCGCAGGGGCCCACCATCAGC 2555
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Db   2501 A----------------------------GTTCCTGTTCTAGTGATGCCACAGTCAGT 2530

Qy   2556 TACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCGGCATGCGCTATCTGGCCACA 2615
          ||| | |  |||  |  | ||||| ||||||| ||||| || |||   || ||  | |
Db   2531 TACGCCAACCTGAAGTTTATGGCAACCCAGATTGCCTCTGGTATGAAGTACCTTTCGTCT 2590

Qy   2616 CTCAACTTTGTACATCGGGACCTGGCCACGCGGAACTGCCTAGTTGGGGAAAATTTCACC 2675
          |||||||||||  || || ||  |||||||| ||  |||||  |||| ||  | |||| ||||
Db   2591 CTCAACTTTGTCCACCGAGATCTGGCCACACGAAACTGTTTAGTGGGCAAGAATTACACC 2650

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Db   2651 ATCAAGATAGCTGATTTTGGCATGAGCAGAAACCTGTACAGTGGTGATTACTACCGGATC 2710

Qy   2736 CAGGGCCGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGGAGTGCATCCTCATGGGGAAG 2795
          ||||||||||  |||||  |||||  |||||||||  |||||||  |||||  | ||||  ||
Db   2711 CAGGGCCGGGCGGTGCTCCCCATTCGCTGGATGTCCTGGGAAAGCATCTTGCTGGGCAAA 2770

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Db   2771 TTCACCACGGCAAGTGATGTGTGGGCCTTTGGGGTGACTCTGTGGGAGACCTTCACCTTT 2830

Qy   2856 TGTAGGGCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGGTCATCGAGAACGCGGGGGAG 2915
          ||   ||  |||||||  |   |||||  | || ||||||||  ||||||||  | || |||
Db   2831 TGCCAGGAGCAGCCCTATTCCCAGCTGTCGGATGAGCAGGTTATCGAGAACACTGGAGAG 2890

Qy   2916 TTCTTCCGGGACCAGGGCCGGCAGGTGTACCTGTCCCGGCCGCCTGCCTGCCCGCAGGGC 2975
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Db   2891 TTCTTCCGAGACCAAGGGAGGCAGATCTATCTCCCTCAACCAGCCCTTTGCCCCGACTCT 2950

Qy   2976 CTATATGAGCTGATGCTTCGGTGCTGGAGCCGGGAGTCTGAGCAGCGACCACCCTTTTCC 3035
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Db   2951 GTGTATAAGCTGATGCTCAGCTGCTGGAGAAGAGAAACCAAGCACCGGCCATCCTTCCAG 3010

Qy   3036 CAGCTGCATCGGTTCCTGGCAGAGGATGCACTCAACACGGTGTGAATCA 3084
          |  | || |   | ||     || | | |  | |        || ||||
Db   3011 GAAATACACCTCCTGCTTCTTCAGCAAGGAGCCGAGTGATGATGCATCA 3059
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RESULT  13
US-08-237-401A-19
; Sequence 19, Application US/08237401A
; Patent No. 5837448
;  GENERAL INFORMATION:
;    APPLICANT:  Lemke Ph.D. et al., Greg E.
;    TITLE OF INVENTION:  PROTEIN-TYROSINE KINASE GENES
;    NUMBER OF SEQUENCES:  54
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Fish & Richardson P.C.
;      STREET:  4225 Executive Square, Suite 1400
;      CITY:  La Jolla
;      STATE:  CA
;      COUNTRY:  US
;      ZIP:  92037
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.25
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/237,401A
;      FILING DATE:  02-MAY-1994
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 07/884,486
;      FILING DATE:  15-MAY-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Haile Ph.D., Lisa A.
;      REGISTRATION NUMBER:  38,347
;      REFERENCE/DOCKET NUMBER:  07251/007001
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (619) 678-5070
;      TELEFAX:  (619) 678-5099
;  INFORMATION FOR SEQ ID NO:  19:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3120 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA
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; IMMEDIATE SOURCE:
; CLONE: Tyro-10
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 485..3047
US-08-237-401A-19

Query Match 16.1%; Score 639.8; DB 2; Length 3120;
Best Local Similarity 56.5%; Pred. No. 4.5e-142;
Matches 1552; Conservative 0; Mismatches 987; Indels 210; Gaps 11;

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Qy    348 CTGCTGCTGCTCTTGGTGGCAAGTGGAGATGCTGACATGAAGGGACATTTTGATCCTGCC 407
Db    509 CTGGTGCTGCTCCTGCTCTTGCTCATCCTGGGTTCTGCAAAAGCTCAGGTTAATCCAGCC 568

Qy    408 AAGTGCCGCTATGCCCTGGGCATGCAGGACCGGACCATCCCAGACAGTGACATCTCTGCT 467
Db    569 ATATGCCGCTATCCTCTGGGCATGTCAGGAGGCCACATTCCAGATGAGGACATCACAGCC 628

Qy    468 TCCAGCTCCTGGTCAGATTCCACTGCCGCCCGCCACAGCAGGTTGGAGAGCAGTGACGGG 527
Db    629 TCAAGTCAGTGGTCAGAATCCACGGCTGCCAAATATGGGAGGCTGGACTCTGAAGAAGGA 688

Qy    528 GATGGGGCCTGGTGCCCCGCAGGGTCGGTGTTTCCCAAGGA---GGAGGAGTACTTGCAG 584
Db    689 GATGGAGCCTGGTGTCCTGAGATTCCAGTGCAACCCGATGACCTGAAGGAATTTCTGCAG 748

Qy    585 GTGGATCTACAACGACTCCACCTGGTGGCTCTGGTGGGCACCCAGGGACGGCATGCCGGG 644
Db    749 ATTGACTTGCGAACCCTACACTTTATCACTCTTGTGGGGACCCAGGGGCGCCATGCAGGG 808

Qy    645 GGCCTGGGCAAGGAGTTCTCCCGGAGCTACCGGCTGCGTTACTCCCGGGATGGTCGCCGC 704
Db    809 GGTCATGGCATTGAATTTGCACCCATGTACAAGATCAACTACAGTCGGGATGGCAGTCGC 868

Qy    705 TGGATGGGCTGGAAGGACCGCTGGGGTCAGGAGGTGATCTCAGGCAATGAGGACCCTGAG 764
Db    869 TGGATCTCCTGGCGTAACCGGCATGGGAAGCAGGTGCTTGATGGAAACAGTAACCCTTAT 928

Qy    765 GGAGTGGTGCTGAAGGACCTTGGGCCCCCCATGGTTGCCCGACTGGTTCGCTTCTACCCC 824
Db    929 GATGTATTCCTGAAGGACTTGGAGCCACCCATCGTCGCCAGATTTGTTCGCCTTATCCCA 988

Qy    825 CGGGCTGACCGGGTCATGAGTGTCTGTCTGCGGGTAGAGCTCTATGGCTGCCTCTGGAGG 884
Db    989 GTCACTGACCACTCCATGAACGTGTGCATGAGGGTTGAGCTTTATGGTTGTGTCTGGCTA 1048

Qy    885 GATGGACTCCTGTCTTACACCGCCCCTGTGGGGCAGACAATGTATTTATCTGAGG----- 939
Db   1049 GATGGCTTGGTATCCTACAATGCTCCAGCTGGACAGCAGTTTGTACTCCCTGGAGGCTCC 1108

Qy    940 -CCGTGTACCTCAACGACTCCACCTATGACGGACATACCGTGGGCGGACTGCAGTATGGG 998
Db   1109 ATCATTTATCTGAATGATTCTGTCTATGATGGAGCTGTTGGGTACAGCATGACTGAAGGG 1168

Qy    999 GGTCTGGGCCAGCTGGCAGATGGTGTGGTGGGGCTGGATGACTTTAGGAAGAGTCAGGAG 1058
Db   1169 ---CTAGGCCAGTTGACTGATGGAGTATCCGGCCTGGATGATTTTACCCAGACCCATGAA 1225

Qy   1059 CTGCGGGTCTGGCCAGGCTATGACTATGTGGGATGGAGCAACCACAGCTTCTCCAGTGGC 1118
Db   1226 TACCACGTGTGGCCTGGCTATGACTACGTGGGATGGCGGAACGAAAGTGCTACCAACGGT 1285

Qy   1119 TATGTGGAGATGGAGTTTGAGTTTGACCGGCTGAGGGCCTTCCAGGCTATGCAGGTCCAC 1178
Db   1286 TTCATTGAGATCATGTTTGAATTTGACCGAATCAGGAATTTTACTACCATGAAGGTCCAC 1345

Qy   1179 TGTAACAACATGCACACGCTGGGAGCCCGTCTGCCTGGCGGGGTGGAATGTCGCTTCCGG 1238
Db   1346 TGCAACAACATGTTTGCTAAAGGTGTGAAGATTTTTAAGGAGGTCCAGTGCTACTTTCGC 1405

Qy   1239 CGTGGCCCTGCCATGGCCTGGGAGGGGGAGCCCATGCGCCACAACCTAGGGGGCAACCTG 1298

Db   1406 TCGG---AAGCCAGCGAGTGGGAACCCACTGCTGTCTACTTTCCCCTGGTCCTGGACGAT 1462

Qy   1299 GGGGACCCCAGAGCCCGGGCTGTCTCAGTGCCCCTTGGCGGCCGTGTGGCTCGCTTTCTG 1358
Db   1463 GTGAACCCCAGTGCCCGGTTTGTCACGGTGCCCCTCCACCACCGAATGGCCAGTGCCATC 1522

Qy   1359 CAGTGCCGCTTCCTCTTTGCGGGGCCCTGGTTACTCTTCAGCGAAATCTCCTTCATCTCT 1418
Db   1523 AAGTGCCAATACCATTTTGCCGACACGTGGATGATGTTCAGCGAGATCACTTTCCAATCA 1582

Qy   1419 GATGTGGTGAACAATTCCTCTCCGGCACTGGGAGGCACCTTCCCGCCAGCCCCCTGGTGG 1478
Db   1583 GATGCTGCAATGTATAACAACTCTGGAGCCCTTCCCACCTCTCCTA---------------- 1628

Qy   1479 CCGCCTGGCCCACCTCCCACCAACTTCAGCAGCTTGGAGCTGGAGCCCAGAGGCCAGCAG 1538
Db   1629 ----------------------------------------TGGCACCCACCACCTATGAT 1648

Qy   1539 CCCGTGGCCAAGGCCGAGGGGAGCCCGACCGCCATCCTCATCGGCTGCCTGGTGGCCATC 1598
Db   1649 CCCATGCTTAAAGTTGATGATAGCAACACTCGGATCCTGATTGGTTGCTTGGTGGCCATC 1708

Qy   1599 ATCCTGCTCCTGCTGCTCATCATTGCCCTCATGCTCTGGCGGCTGCACTGGCGCAGGCTC 1658
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Qy   1659 CTCAGCAAGGCTGAACGGAGGGTGTTGGAAGAGGAGCTGACGGTTCACCTCTCTGTCCCT 1718
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Qy   1719 GGGGACACTATCCTCATCAACAACCGCCCAGGTCCTAGAGAGCCACCCCCGTACCAGGAG 1778
Db   1829 AGCGAGTCCAGCATGTTCAATAACAACCGCTCCTC------ATCACCAAGTGAACAGGAG 1882

Qy   1779 CCCCGGCCTCGTGGGAATCCGCCCCACTCCGCTCCCTGTGTCCCCAATGGCTCTGCGTTG 1838
Db   1883 TCCAACTCTACTTATGATCGAATCTTCCCCCTTCGCCCTGACTACCAGGAGCCATCCAGA 1942

Qy   1839 CTGCTCTCCAATCCAGCCTACCGCCTCCTTCTGGCCACTTACGCCCGTCCCCCTCGAGGC 1898
Db   1943 CTGATC---------------------------------------------------- 1948

Qy   1899 CCGGGCCCCCCCACACCCGCCTGGGCCAAACCCACCAACACCCAGGCCTACAGTGGGGAC 1958
Db   1949 ------------------------CGAAAGCTTCCAGAGTTTGCTCCAGGAGAGGAGGA 1983

Qy   1959 TATATGGAGCCTGAGAAGCCAGGCGCCCCGCTTCTGCCCCCACCTCCCCAGAACAGCGTC 2018
Db   1984 GTCAGGGTGCAGTGGTGTTGTGAAGCCGGCC-----CAGCCCAATGGACCTGAGGGCGTG 2038

Qy   2019 CCCCATTATGCCGAGGCTGACATTGTTACCCTGCAGGGCGTCACCGGGGGCAACACCTAT 2078
Db   2039 CCCCACTATGCAGAAGCCGACATAGTGAATCTCCAGGGAGTGACAGGTGGCAACACCTAC 2098

Qy   2079 GCTGTGCCTGCACTGCCCCCAG---GGGCAGTCGGGGATGGGCCCCCCAGAGTGGATTTC 2135
Db   2099 TGTGTGCCTGCTGTAACCATGGATCTGCTATCGGGGAAAGATGTGGCTGTGGAAGAGTTC 2158

Qy   2136 CCTCGATCTCGACTCCGCTTCAAGGAGAAGCTTGGCGAGGGCCAGTTTGGGGAGGTGCAC 2195
Db   2159 CCCAGGAAACTGTTGGCCTTCAAGGAGAAGCTGGGAGAAGGCCAGTTTGGGGAGGTTCAT 2218

Qy   2196 CTGTGTGAGGTCGACAGCCCTCAAGATCTGGTCAGTCTTGATTTCCCCCTTAATGTGCGT 2255
Db   2219 CTCTGTGAAGTGGAGGGAATGGAAAAATTCAAAGACAAAGATTTTGCACTAGATGTCAGT 2278

Qy   2256 AAGGGACACCCTTTGCTGGTAGCTGTCAAGATCTTACGGCCAGATGCCACCAAGAATGCC 2315
Db   2279 GCCAACCAGCCTGTCCTGGTGGCCGTGAAAATGCTCCGAGCAGATGCCAACAAGAATG-- 2336

Qy   2316 AGCTTCTCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGGTGAAGATCATGTCGAGGCTC 2375
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Db    2337 ----------------CCAGGAATGATTTTCTTAAGGAGATCAAGATCATGTCTCGGCTC 2380

Qy    2376 AAGGACCCCAACATCATTCGGCTGCTGGGCGTGTGTGTGCAGGACGACCCCCTCTGCATG 2435

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Qy    2496 GACAAGGCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTGCGCAGGGGCCCACCATCAGC 2555

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Qy    2556 TACCCAATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCGGCATGCGCTATCTGGCCACA 2615

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Db    3011 GAAATACACCTCCTGCTTCTTCAGCAAGGAGCCGAGTGATGATGCATCA 3059


RESULT  14
US-08-286-305A-4
; Sequence 4, Application US/08286305A
; Patent No. 5766863
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Sadick, Michael D.
;    APPLICANT:  Shelton, David L.
;    APPLICANT:  Wong, Wai Lee Tan
;    TITLE OF INVENTION:  KINASE RECEPTOR ACTIVATION ASSAY
;    NUMBER OF SEQUENCES:  11
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  5.25 inch, 360 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/286,305A
;      FILING DATE:  05-AUG-1994
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/170558
;      FILING DATE:  20-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Lee, Wendy M.
;      REGISTRATION NUMBER:  00,000
;      REFERENCE/DOCKET NUMBER:  854C1P1
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1994
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  4:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2820 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  linear
US-08-286-305A-4


  Query Match             4.6%;  Score 182.2;  DB 1;  Length 2820;
  Best Local Similarity   59.9%;  Pred. No. 7.9e-34;
  Matches  348;  Conservative    0;  Mismatches  218;  Indels   15;  Gaps    2;

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Qy    2382 CCCAACATCATTCGGCTGCTGGGCGTGTGTGTGCAGGACGACCCCCTCTGCATGATTACT 2441

Db    2054 CAGCACATCGTGCGCTTCTTCGGCGTCTGCACCGAGGGCCGCCCCCTGCTCATGGTCTTT 2113

Qy    2442 GACTACATGGAGAACGGCGACCTCAACCAGTTCCTCAGTGCCCACCAGCTGGAGGACAAG 2501

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Qy    2502 GCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTGCGCAGGGGCCCACCATCAGCTACCCA 2561

Db    2173 ----------GCTGCTGGCTGGTGGGGAGGATGTGGCTCCAGGCCCCCTGGGTCTGGGG 2221

Qy    2562 ATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCGGCATGCGCTATCTGGCCACACTCAAC 2621

Db    2222 CAGCTGCTGGCCGTGGCTAGCCAGGTCGCTGCGGGGATGGTGTACCTGGCGGGTCTGCAT 2281

Qy    2622 TTTGTACATCGGGACCTGGCCACGCGGAACTGCCTAGTTGGGGAAAATTTCACCATCAAA 2681

Db    2282 TTTGTGCACCGGGACCTGGCCACACGCAACTGTCTAGTGGGCCAGGGACTGGTGGTCAAG 2341

Qy    2682 ATCGCAGACTTTGGCATGAGCCGGAACCTCTATGCTGGGGACTATTACCGTGTGCAGGGC 2741

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Db    2402 CGCACCATGCTGCCCATTCGCTGGATGCCGCCCGAGAGCATCCTGTACCGTAAGTTCACC 2461

Qy    2802 ACTGCGAGTGACGTGTGGGCCTTTGGTGTGACCCTGTGGGAGGTGCTGATGCTCTGTAGG 2861

Db    2462 ACCGAGAGCGACGTGTGGAGCTTCGGCGTGGTGCTCTGGGAGATCTTCACCTACGGCAAG 2521

Qy    2862 GCCCAGCCCTTTGGGCAGCTCACCGACGAGCAGGTCATCGA 2902

Db    2522 ---CAGCCCTGGTACCAGCTCTCCAACACGGAGGCAATCGA 2559
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RESULT 15
US-08-441-104A-4
; Sequence 4, Application US/08441104A
; Patent No. 5891650
;  GENERAL INFORMATION:
;    APPLICANT:  Godowski, Paul J.
;    APPLICANT:  Mark, Melanie R.
;    APPLICANT:  Sadick, Michael D.
;    APPLICANT:  Shelton, David L.
;    APPLICANT:  Wong, Wai Lee Tan
;    TITLE OF INVENTION:  KINASE RECEPTOR ACTIVATION ASSAY
;    NUMBER OF SEQUENCES:  11
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Genentech, Inc.
;      STREET:  460 Point San Bruno Blvd
;      CITY:  South San Francisco
;      STATE:  California
;      COUNTRY:  USA
;      ZIP:  94080
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  3.5 inch, 720 Kb floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  patin (Genentech)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/441,104A
;      FILING DATE:  15-MAY-1995
;      CLASSIFICATION:  435
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/286305
;      FILING DATE:  05-AUG-1994
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/170558
;      FILING DATE:  20-DEC-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  08/157563
;      FILING DATE:  23-NOV-1993
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Lee, Wendy M.
;      REGISTRATION NUMBER:  00,000
;      REFERENCE/DOCKET NUMBER:  854C1P1C2
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  415/225-1994
;      TELEFAX:  415/952-9881
;      TELEX:  910/371-7168
;  INFORMATION FOR SEQ ID NO:  4:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2820 bases
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  linear
US-08-441-104A-4
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Query Match 4.6%; Score 182.2; DB 2; Length 2820;
Best Local Similarity 59.9%; Pred. No. 7.9e-34;
Matches 348; Conservative 0; Mismatches 218; Indels 15; Gaps 2;

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Qy   2322 TCCTTGTTCTCCAGGAATGATTTCCTGAAAGAGGTGAAGATCATGTCGAGGCTCAAGGAC 2381
          |||  |    |  || | || ||||     ||||   || |  |  | | |||  || ||
Db   1994 TCCGAGAGTGCTCGGCAGGACTTCCAACGTGAGGCTGAGCTGCTCACCATGCTGCAGCAC 2053

Qy   2382 CCCAACATCATTCGGCTGCTGGGCGTGTGTGTGCAGGACGACCCCCTCTGCATGATTACT 2441
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Db   2054 CAGCACATCGTGCGCTTCTTCGGCGTCTGCACCGAGGGCCGCCCCCTGCTCATGGTCTTT 2113

Qy   2442 GACTACATGGAGAACGGCGACCTCAACCAGTTCCTCAGTGCCCACCAGCTGGAGGACAAG 2501
          || || |||  | |||| ||||||||||  |||||| |  ||||    |  || | |||
Db   2114 GAGTATATGCGGCACGGGGACCTCAACCGCTTCCTCCGATCCCATGGACCTGATGCCAA- 2172

Qy   2502 GCAGCCGAGGGGGCCCCTGGGGACGGGCAGGCTGCGCAGGGGCCCACCATCAGCTACCCA 2561
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Db   2173 ----------GCTGCTGGCTGGTGGGGAGGATGTGGCTCCAGGCCCCCTGGGTCTGGGG 2221

Qy   2562 ATGCTGCTGCATGTGGCAGCCCAGATCGCCTCCGGCATGCGCTATCTGGCCACACTCAAC 2621
          |||||||   |||||   |||| |||| | || |||   || |||||    ||  |
Db   2222 CAGCTGCTGGCCGTGGCTAGCCAGGTCGCTGCGGGGATGGTGTACCTGGCGGGTCTGCAT 2281

Qy   2622 TTTGTACATCGGGACCTGGCCACGCGGAACTGCCTAGTTGGGGAAAATTTCACCATCAAA 2681
          ||||| || ||||||||||||||| || ||||| ||||| ||  |     |     ||||
Db   2282 TTTGTGCACCGGGACCTGGCCACACGCAACTGTCTAGTGGGCCAGGGACTGGTGGTCAAG 2341

Qy   2682 ATCGCAGACTTTGGCATGAGCCGGAACCTCTATGCTGGGGACTATTACCGTGTGCAGGGC 2741
          || |  || ||||||||||||| || |  ||||        ||||||||||||||    |||
Db   2342 ATTGGTGATTTTGGCATGAGCAGGGATATCTACAGCACCGACTATTACCGTGTGGGAGGC 2401

Qy   2742 CGGGCAGTGCTGCCCATCCGCTGGATGGCCTGGGAGTGCATCCTCATGGGGAAGTTCACG 2801
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Db   2402 CGCACCATGCTGCCCATTCGCTGGATGCCGCCCGAGAGCATCCTGTACCGTAAGTTCACC 2461

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Search completed: December 8, 2001, 21:00:30
Job time: 4924 sec